STIC-Biot ch/ChemLib

From: Sent:

Schultz, James

Wednesday, May 07, 2003 9:22 AM STIC-Biotech/ChemLib

To: Subject:

Sequence search 10/057,550

MAY -7 2003

HECK/CHEM. CIVISION (STIC) Whoops, sorry, the corrected file name is now listed in the title above. .

-----Original Message-----

From:

STIC-Biotech/ChemLib

Sent:

Wednesday, May 07, 2003 9:06 AM

To:

Schultz, James

Subject:

RE: Sequence search pct/us02/04123

This Application Number does not have a valid CRF - please provide the necessary correction or another application

maude

-----Original Message-----

From:

Schultz, James

Sent:

Wednesday, May 07, 2003 8:36 AM

To:

STIC-Biotech/ChemLib

Subject: Sequence search pct/us02/04123

Hello,

I need a length limited nucleotide sequence search performed on SEQ ID NO: 64 in the above entitled application, where the maximum size of the returned hit is no more than 50 nucleotides long. Thanks, Doug Schultz

J. Douglas Schultz, Ph.D. AU 1635 (Biotechnology) Patent Examiner United States Patent and Trademark Office (703) 308-9355 (703) 746-3973 (fax) Office: CM1 12E18 Mail: CM1 11E12

Edward Hart Technicai Info. Specialist STIC/Biotech CMI 6B02 Tel: 305-9203

Searcher:
Phone:
Location:
Date Picked Up: 5/8/1/25
Date Completed: 5/17/05
Searcher Prep/Review:
Clerical:
Online time:

TYPE OF SEARCH;
NA Sequences:
AA Sequences:
Structures:
Bibliographic:
Litigation:
Full text:
Patent Family:
Other:

VENDOR/COST (where applic.)
STN:
DIALOG:
Questel/Orbit:
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Lexis/Nexis:
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Other (specify):

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STIC SEARCH RESULTS

Biotech-Chem Library

Questions about the scope or the results of the search? Contact the searcher or contact:

Mary Hale, Information Branch Supervisor 308-4258, CM1-1E01

V	antary Results / ccaback. Oill
>	I am an examiner in Workgroup: Example: 1610
>	Relevant prior art found, search results used as follows:
	☐ 102 rejection
	☐ 103 rejection
	Cited as being of interest.
	Helped examiner better understand the invention.
	Helped examiner better understand the state of the art in their technology.
	Types of relevant prior art found:
	Foreign Patent(s)
	Non-Patent Literature (journal articles, conference proceedings, new product announcements etc.)
>	Relevant prior art not found:
	Results verified the lack of relevant prior art (helped determine patentability).
	Results were not useful in determining patentability or understanding the invention.
Co	mments:

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score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result ĕ Match AR100923 AR100938 A63567 AR142002 AR170583 AX164948 AR079693 AR081223 AR110776 AX157157 AR079693 AR197894 AR090860 AR197895 ARI10776 Sequence AR177157 Sequence AR079693 Sequence AR079693 Sequence AR142002 Sequence AR14002 Sequence AR110053 Sequence AR1109540 Sequence AR100937 Sequence AR100937 Sequence AR100937 Sequence AR100937 Sequence AR100938 Sequence AR100938 Sequence AR10938 Sequence AR10958 Sequence AR10959 Sequence AR10958 Sequence AR10958 Sequence AR10958 Sequence AR10958 Sequence AR15572 Sequence AR16573 Sequence AR16773 Sequence AR16773 Sequence AR16773 Sequence AR16773 Sequence AR16773 Sequence AR16773 Sequence AX248756 Sequence AR090859 Sequence AR197894 Sequence AR090860 Sequence AR197895 Sequence 196168 Sequence 5 196169 Sequence 5 AX183838 Sequence L27066 Human neuro AX287576 Sequence AX287580 Sequence AX277277 Sequence L58003 Sequence 11 A52104 Sequence 7 A52105 Sequence 7 A52132 Sequence 5 Description

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REFERENCE AUTHORS		ORGANISM	SOURCE	SEGMENT	KEYWORDS	VERSION	ACCESSION ·	DEFINITION	LOCUS	HUMRAF12	RESULT 1
<pre>1 (bases 1 to 49) Bonner,T.I., Kerby,S.B., Sutrave,P., Gunnell,M.A., Mark,G. and Rapp,U.R.</pre>	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.	Homo sapiens	Homo sapiens (tissue library: of Lawn et al.) DNA.	2 of 9	<pre>c-myc proto-oncogene; raf protein; raf proto-oncogene.</pre>	L00213.1 GI:190830	L00213 ·	Human c-raf-1 proto-oncogene, exon 2, clones lambda-(2,13).	HUMRAF12 49 bp DNA linear PRI 08-JAN-1995		

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Patent: WO 0166800-A 835 13-SEP-2001;
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH
                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata;
Mammalia; Eutheria; Primates; Catarrhini
1 (bases 1 to 31)
Cargill,M., Ireland,J.S. and Lander,E.S.
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Sequence 835 from Patent W00166800
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AR090860
                                                    1 (bases 1 to 26)
Chenchik, A., Jokhadze, G. and Bibilashvilli, R.
Methods of assaying differential expression
Patent: US 5994076-A 980 30-NOV-1999;
                                                                                                               Unknown
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AR197894
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28; Conservative
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Patent: US 6352829-A 979 05-MAR-2002;
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Chenchik, A., Jokha
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1 (bases 1 to 28)
Chenchik, A., Jokhadze,
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Unclassified.
Unclassified.
1 (bases 1 to 26)
1 (bases 1, Jokhadze,G. and Bibilashvilli,R. Chenchik,A., Jokhadze,G. and Bibilashvilli,R. Methods of assaying differential expression Patent: US 6352829-A 980 05-MAR-2002;
Location/Qualifiers
  Sequence
196169
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196168
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AR197895
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1 (bases 1 to 26)

Calabretta, B. and Skorski, T.

Antisense oligonucleotides targeting cooperating oncogenes

Patent: US 5734039-A 5 31-MAR-1998;
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l Similarity 100.0%;
26; Conservative
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26; Conser
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4 c 14 g
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AX157157
AX157157.1 G
                            Shimkets,R.A. and Leach,M.
Nucleic acids containing single nucle
methods of use thereof
Patent: WO 0140521-A 485 07-JUN-2001;
Curagen Corporation (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Calabretta, B. and Skorski, T. Antisense oligonucleotides targetin Patent: US 5734039-A 6 31-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                 Kasid, U., Gokhale, P., Dritschilo, A. a Liposomes containing oligonucleotides Patent: US 6126965-A 2 03-OCT-2000;
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AR110776
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Catarrhini; Hominidae
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PAT 14-FEB-2001

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                        28;
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Patients, G.N. and Felber, B.K.
Nucleic acid constructs containing H
inhibitory/instability regions and mm
Patent: US 5972596-A 17 26-OCT-1999;
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l Similarity 73.7%;
28; Conservative
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Method of eliminating inhibitory/ instability
Patent: US 5965726-A 17 12-OCT-1999;
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Unclassified.
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25. .26
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Pred. No. 1.5e+06;
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Pred. No. 1.3e-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     QΥ
                                                                                                                                                                                                                                                                                                                                                                                 Locus
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AR142002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       KEYWORDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /ERSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CUS
                       ORGANISM
                                                                                                                                                                                    Query Match
Best Local
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                                                                                                                                             2207 TCACAGGGCCTTAACTTCATGTTGCCTTCTTTTCTATC
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                                                                                                                                  2 TCTCAGATACCTAGCTTCATATTGCCTTATTGTCTACC 39
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                                                  AX164948
Sequence 143
AX164948
AX164948.1
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Sequence
AR170583
  Eukaryota; Metazoa;
Mammalia; Eutheria;
                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AR142002
Sequence
AR142002
                                human
                                                                                                                                                                                                                                                                1 (bases 1 to 43)
Pavlakis,G.N. and Felber,B.K.
Method of eliminating inhibitory/instability
Patent: US 6291664-A 17 18-SEP-2001;
                                                                                                                                                                                                                                                                                                                     Unknown
                                                                                                                                                                                                                                                                                                                                Unknown.
                                                                                                                                                                                                                                                                                                                                                   AR170583.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity 28; Conserv
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Pavlakis,G.N. and Felber,B.K.
Method of eliminating inhibitory/instability
Patent: US 6174666-A 17 16-JAN-2001;
                                                                                                                                                                                                                                                                                                            Unclassified.
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11 c 5 q
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11 c 5 g
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                                                    GI:14545777
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                                                                                                                                                                                                                                                                                                                                                                   43 bp
from patent US 6291664.
                                                                                                                                                                                    0.7%;
73.7%;
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                                                                       Patent
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 Chordata;
Primates;
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                                                                                                                                                                          Score 22; DB Pred. No. 1.5e 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 22; DB (
Pred. No. 1.5e+
0; Mismatches
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                                                                      50 bp
WO0138586.
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s 6174666.
Craniata; V
Catarrhini;
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1.5e+06;
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          Vertebrata;
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         Euteleostomi;
                                                                               PAT 22-JUN-2001
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                                                                                                                                                                                                                                                                             of mRNA
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                                                                                                                                                                         0;
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REFERENCE 1 (bases 1 to 50)
ANTHORS Shinkets, R.A. and Leach, M.
TITLE Nucleic acids containing single nucleotide polymorphisms and methods of use thereof
JOURNAL Patent: WO 0138586-A 143 31-MAY-2001;
Curagen Corporation (US)
FEATURES Location/Qualifiers
Source //OB_XTEF="taxon:9606"

misc_feature 25..26
//OB_XTEF="taxon:9606"
//OB_XTEF="Nucleotide deleted between bases 25 and 26
Accession number cg43945210"
variation 26
Accession number cg43945210"

variation 9 a 11 c 8 g 22 t
ORIGIN

Query Match 0.7%; Score 22; DB 6; Length 50;
Best Local Similarity 83.3%; Pred. No. 1.5e+06;
Matches 25; Conservative 0; Mismatches 5; Indels 0; Gaps Qy 2804 AAACAAACAAACAACCCTTGTTTTCTT 2833
Qy 2804 AAACAAACAAACAAACAACCCTTGTTTTCTT 2833
Db 33 AAACAAACAAACAAACAACCTTGTGTTCTGT 4

Search completed: May 16, 2003, 09:18:22
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Title:
Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq
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2977
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GenCore version 5.1.4_p5_4578 Copyright (c) 1993. - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                            /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1980.DAT:
                                                                                                                                                                                                                                                                                                                                                                                                                 'SIDS2/gcgdata/geneseq
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AAA07027
AAA3030347
AAI30349
AAK66891
AAV20480
ABK66892
AAL31396
AAH444455
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Raf-1
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21	21	21	21	21	21.2	21.4	21.4	21.4	21.4	21.4	21.4	21.6	22	22	22	22	22	22	22	22.2	22.2	22.4	22.4	22.4	22.4	23	23	23	23	23	23	24	24	24	24.2
0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7					0.7	0.7	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	٠
40	27	27	27	21	35	40	40	40	40	40	40	50	50	47	43	41	30	30	30	50	27	50	27	27	27	23	23	23.	23	23	23	ω ω	31	24	35
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AAV05431	ABL50208	ABL50176	AAH41589	AAF97098	AAT93823	AAA55866 .	AAA55851	AAA72893	AAA72878	AAV05418	AAV05430	AAI75927	ABL00152	AAZ69520	AAQ50213	AAT93841	ABL50209	ABL50177	AAH41590	AAI73544	AAT93834	AAL31673	ABL50229	ABL50197	AAH41610	AAA26246	AAA26245	AAA26244	AAA26243	AAA26242	AAA26239	AAA29998	AAA30000	AAV33096	AAA29996
					4.																								•				•		
iA us	PCR	Human c-Rafl PCR p	Human c-Rafl PCR p	Human gene single	Antitumoural phosp	GAD plasmid RAS in	RAF PCR primer SEQ	Yeast PCR primer R	gene	Primer used in pro	S us	Human silent SNP c	Human silent nonco	Human map-related	c-fos position 339	Phosphodiester oli	Raf1 PCR	PCR	PCR	Human silent SNP c	Phosphodiester oli	SNP oligonu	c-Rafl PCR	c-Rafl PCR	an c-Rafl PCR	af antisense o	af antisense o	af antisense	af antisense o	antisense	C-raf antisense ol		in hybridize		Hairpin hybridizer

ALIGNMENTS

AAA07016 standard; DNA; 42 BP

AAA07016;

03-JUL-2000 (first entry)

Raf-1 mutagenic PCR primer, SEQ ID NO:13.

Raf-1; CAP kinase; phosphorylation; ceramide-activated protein kinase; mutagenesis; lipopolysaccharide; LPS; endotoxin; sphingomyelin PCR primer;

Mammalia.

US6040149-A

21-MAR-2000

10-JAN-1997; 97US-0785247.

11-JAN-1996; 96US-0009900.

(SLOK) SLOAN KETTERING INST CANCER RES

Liu J, Kolesnick RN;

WPI; 2000-270133/23.

latexin

Novel method of identifying agents capable of inhibiting

Novel method of identifying agents capable of inhibiting lipopolysaccharide induced threonine phosphorylation by ceramide-activated protein kinase

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RESULT 2
AAA07027
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Best Local S
Matches 38
   WPI;
                          Zhang Y, Liu J, Kolesnick
                                                    (SLOK ) SLOAN KETTERING INST CANCER RES
                                                                                          11-JAN-1996;
                                                                                                                     10-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   to ceramide. The ceramide in turn stimulates the kinase to phosphorylate protein substrates which can then mediate signal transduction. The CAP kinase is also stimulated by the bacterial endotoxin lipopolysaccharide (LPS), which is thought to mimic the second messenger function of ceramide. The methods are useful for identifying agents that inhibit lipopolysaccharide-induced Thr phosphorylation by CAP kinase. The agents identified using the method are useful for treating disorders associated with aberrant phosphorylation of target molecules by CAP kinase, e.g., inflammatory disorders (such as rheumatoid arthritis), ulcerative collitis, graft versus host disease, lupus
                                                                                                                                                    21-MAR-2000
                                                                                                                                                                                  US6040149-A
                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                Raf-1; CAP kinase; phosphorylation; ceramide-activated protein
                                                                                                                                                                                                                                                                                                                  Raf-1 mutagenic PCR primer,
                                                                                                                                                                                                                                                                                                                                                                                AAA07027;
                                                                                                                                                                                                                                Mammalia.
                                                                                                                                                                                                                                                           sphingomyelin
                                                                                                                                                                                                                                                                            mutagenesis;
                                                                                                                                                                                                                                                                                                                                                     03-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                            AAA07027 standard; DNA; 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              erythematosus, HIV, infection, disorders associated with poor streamouth, and septic shock. Sequences AAA07016-A07019 and AAA07027 represent PCR primers used in an exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           weight of 100-110 kD. It is an upstream participant in a sphingomyelin signal transduction pathway which uses ceramide as a second messenger. This pathway is initiated by tumour necrosis factor-alpha (TNF-alpha) and interleukin-beta (IL-beta), causing the hydrolysis of sphingomyelin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a novel method of determining whether an agent is capable of specifically inhibiting the ability of a ceramide-activated protein (CAP) kinase to phosphorylate the threonine residue in a polypeptide containing a Thr-Pro- or Thr-Leu-Pro motif. In
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Raf-1, epidermal growth factor receptor (EGFR), or suitable fragments thereof. The CAP kinase is membrane bound and has an apparent molecular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              particular,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example VI; Column
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  lipopolysaccharide ceramide-activated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      generate mutant
2000-270133/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 9 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the peptide substrate that is specifically phosphorylated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                        signal transduction pathway;
                                                                                      96US-0009900
                                                                                                                     97US-0785247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Flag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                56; 84pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  induced threonine phosphorylation by protein kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      peptide/Raf-1
                                                                                                                                                                                                                                                                                                                                                                                                            ВP
                            RN
                                                                                                                                                                                                                                                                                                                    SEQ ID NO: 27:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 35.6;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                         LPS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequences
                                                                                                                                                                                                                                                        PCR primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  via overlap extension
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                   kinase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              stem cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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cc weight of 100-110 kD. It is an upstream participant in a sphingomyelin consignal transduction pathway which uses ceramide as a second messenger. Cc This pathway is initiated by tumour necrosis factor-alpha (TNF-alpha) cc and interleukin-beta (II-beta), causing the hydrolysis of sphingomyelin cc to ceramide. The ceramide in turn stimulates the kinase to c phosphorylate protein substrates which can then mediate signal cc transduction. The CAP kinase is also stimulated by the bacterial cc medical lipopolysaccharide (LPS), which is thought to mimic the second conditions that inhibit lipopolysaccharide-induced. The methods are useful for identifying agents that inhibit lipopolysaccharide of using the method are useful for creating disorders associated with aberrant phosphorylation by creating disorders associated with aberrant phosphorylation of target conclecules by CAP kinase, e.g., inflammatory disorders (such as rheumatoid arthritis), ulcerative colliis, graft versus host disease, lupus crythematosus, HIV, infection, disorders associated with poor stem cell crythematosus, HIV, infection, disorders associated with poor stem cell cromber in the collician and exemplification of the present invention to generate mutant Flag peptide/Raf-1 sequences via overlap extension
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             residue in a polypeptide containing a Thr-Pro- or Thr-Leu-Pro motif. In particular, the peptide substrate that is specifically phosphorylated is Raf-1, epidermal growth factor receptor (EGFR), or suitable fragments
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a novel method of determine capable of specifically inhibiting the ability
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ceramide-activated protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Raf-1, epidermal growth factor receptor (EGFR), or suitable fragments thereof. The CAP kinase is membrane bound and has an apparent molecular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example VI; Column 56; 84pp; English
                                                            Flag peptide/Raf-1 sequences via
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CAP) kinase to phosphorylate the threonine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of determining whether an agent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    of
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В QΥ Best Local Query Match 919 1 CACATGGTCAGCACCGTGCTACCGGTGGACAGCAGGATG CACATGGTCAGCACCACGCTGCCTGTGGACAGCAGGATG 957 35; Similarity Conservative 1.1%; 13 G; 6 T; 0 other; 0 Score 32.6; Pred. No. 1: Mismatches DВ 39 21; 4 Length Indels 39; 0 Gaps

0

Sequence

39 BP;

9

A,

11 C;

BXXXX AAA07017 standard; DNA; 42

AAA07017;

03-JUL-2000 (first entry)

Raf-1 mutagenic PCR primer, SEQ ID NO:14.

Raf-1; CAP kinase; phosphorylation; ceramide-activated protein kinase; mutagenesis; lipopolysaccharide; LPS; endotoxin; sphingomyelin signal transduction pathway; PCR primer; ss.

Synthetic. Mammalia.

US6040149-A

21-MAR-2000

11-JAN-1996; 10-JAN-1997; 96US-0009900 97US-0785247

(SLOK) SLOAN KETTERING INST CANCER RES

Zhang Y, Liu Ţ, Kolesnick

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RESULT 4
AA130347
ID AA13
XX AA12
XX AA12
XX Humm
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a novel method of determining whether an agent clis capable of specifically inhibiting the ability of a ceramide-activated protein (CRP) kinase to phosphorylate the threonine craidle activated protein (CRP) kinase to phosphorylate the threonine residue in a polypeptide containing a Thr-Pro- or Thr-Leu-Pro motif. In CRP kinase is membrane bound and has an apparent molecular craidle transduction pathway which uses ceramide as a second messenger. This pathway is initiated by tumour necrosis factor-alpha (TRP-alpha) craidle. The ceramide in turn stimulates the kinase to phosphorylate protein substrates which can then mediate signal cransduction. The CAP kinase is also stimulated by the bacterial cransduction. The CAP kinase is also stimulated by the bacterial cransduction. The CAP kinase is also stimulated by the bacterial concessorer function of coramide. The mediate signal the second concessorer function of coramide. The mediate signal cransduction. The CAP kinase is also stimulated by the bacterial concessorer function of coramide. The mediate signal the second concessorer function of coramide. The mediate signal concessorer function of coramide.
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                                   07-MAR-2001; 2001WO-US07268
                                                                                                        13-SEP-2001.
                                                                                                                                                                                                                                                                                                              Key
Variation
                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAI30347;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 42 BP; 8 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example VI; Column 56; 84pp; English
                                                                                                                                                                            WO200166800-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human single nucleotide polymorphism (SNP) RAF1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAI30347 standard; DNA; 31 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-270133/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              916 GTCCACATGGTCAGCACCACGCTGCCTGTGGACAGCAGGATG 957
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            generate mutant Flag peptide/Raf-1 sequences via
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GTCCACATGGTCAGCGTCGTGCTACCGGTGGACAGCAGGATG 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   resequence; genotype; disease; forensic; paternity testing; nucleotide polymorphism; SNP; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   l Similarity
36; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                             Location/Qualifiers replace(16,T)
/*tag= a
                                                                                                                                                                                                                                         /standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11 C; 15 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 32.4;
Pred. No. 13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6
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ABK66891
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to the identification of nucleic acid molecules (AAI29513-AAI31314) from the human genome which include polymorphic sites which can predispose individuals to disease. Various genes from a number of individuals were resequenced and single nucleotide polymorphisms (SNPS) in these genes discovered. The method is useful for predicting the presence, absence or severity of a particular phenotype or disorder (e.g. diabetes) associated with a particular genotype. The nucleic acids
                                                        Producing sub-population of labeled nucleic acids, useful for analysing differences in RNA profiles between several different physiological sources, using set of distinct gene specific primers -
                                                                                                            WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2144 CCCTTTCTCCAGAGGCAGAACACATGTTTTC 2174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acid molecules from the human genome which include polymorphic sites, useful in methods for predicting the presence, absence or severity of a particular phenotype or disorder (e.g. diabetes) associated with a particular genotype .
                                                                                                                                    Chenchik A,
                                                                                                                                                                                                                                                                                                                 Primer; ss; DNA microarray; differential expression analysis; human
                                                                                                                                                                                                                                                                                                                                                                   02-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 80; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cargill M,
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22-MAY-2000;
                                                                                                                                                             (CLON-) CLONTECH LAB INC
                                                                                                                                                                                       21-MAY-1997;
                                                                                                                                                                                                               05-JAN-1999;
                                                                                                                                                                                                                                       05-MAR-2002
                                                                                                                                                                                                                                                                US6352829-B1
                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                         Human gene specific PCR primer
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                                                                                                                                                                                                                                                                                                                                                                                                                    ABK66891 standard; DNA; 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             testing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      containing the
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                                                                                                            2002-314699/35.
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                                                                                                                                    Jokhadze G,
                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      polymorphic sites may be
                                                                                                                                                                                       97US-085999E
                                                                                                                                                                                                                99US-0225928
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                                                                                                                                    Bibilashvilli R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5 G;
                                                                                                                                                                                                                                                                                                                                                                                                                    ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 31;
Pred. No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9 T;
                                                                                                                                                                                                                                                                                                                                           #979
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 22;
29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         useful in forensics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 31;
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The invention relates to producing a sub-population of labeled nucleic acids (NAs) comprising contacting a NA sample from a physiclogical

Example 3; SEQ ID No 979;

11pp; English

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RESULT 6
AAV20480/c
ID AAV204
XX AAV204
XX AAV204
XX 17-JUN
AC Human
DT 17-JUN
XX Human;
KW Cancer
XX Synthe
OS Synthe
FF Key
FF Allele
FT allele
FT Allele
FT Allele
FT SEP
XX US5734
XX US5734
XX US5734
XX US5734
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XX US5734
XX Antical
PT Calabr
XX Antical
PT AAV204
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Best Local (
              Anticancer composition comprising two anti-sense oligo:nucleotide(s)
- targetting cytoplasmic and nuclear oncogene(s);
                                                               WPI; 1998-229882/20
                                                                                                                                                                      15-SEP-1994;
                                                                                                                                                                                                        15-SEP-1994;
                                                                                                                                                                                                                                                                           US5734039-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human c-raf antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1704 CAGTITCCAGTCGGATGTCTACTCCTAT 1731
                                                                                                                                     (UYJE-) UNIV JEFFERSON THOMAS
                                                                                                                                                                                                                                           31-MAR-1998
                                                                                                                                                                                                                                                                                                                                               allele
                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                              cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; oncogene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAV20480;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAV20480 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electro format directly from USPTO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               between several different physiological sources, where the method comprises producing subpopulation of labeled NAs for the different physiological sources, comprising the populations for each physiological sources to identify differences in the population, where the comparison is preferably performed by hybridising the labeled NAs for each of the distinct physiological sources to an array of probe NAs stably associated with the surface of a substrate to produce a hybridisation pattern for each of the sources, and comparing the patterns for each of the sources, where differential gene expression assays are utilised in differential expression analysis of diseased a normal tissue e.g. neoplastic a normal tissue, or different tissue or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 subtissue types. The present sequence is a human gene specific PCR primer used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    primer. The method is useful for producing a sub-population of labele NAs which is useful for analysing the differences in the RNA profiles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   http.wipo.seqdata.uspto.gov/sequence.html?DocID=6352829B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       each gene specific primer has a sequence compumnia, and each labeled NA is generated using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                source, with a pool conditions to enzym
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       H
                                                                                                                                                                                                                                                                                                                                                                                             sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CAGTTTCCAGTCGGATGTCTACTCCTAT 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28;
                                                                                                                                                                                                                                                                                                                                                                                                                                              antisense
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                                                                                                     В,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.9%;
Lity 100.0%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           a pool of 50 distinct gene specific primers under suitable enzymatically generate sub-non-lation of the specific primers under suitable
                                                                                                   Skorski
                                                                                                                                                                    94US-0306691
                                                                                                                                                                                                      94US-0306691
                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                             proto-oncogene;
                                                                                                                                                                                                                                                                                                                                                                                                                                         proto-oncogene; neoplastic disease; anticancer;
oligonucleotide; phosphorothicate; ss.
                                                                                                                                                                                                                                                                                                      "phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       has a sequence complementary to a distinct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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Pred. No. 2.2
0; Mismatches
            oncogene(s)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ng a single gene specifi
a sub-population of lab
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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Best Local
     MAS which is useful for analysing the differences in the RNA profiles between several different physiological sources, where the method comprises producing subpopulation of labeled NAS for the different physiological sources, comprising the populations for each physiological sources comprising the populations for each physiological source to identify differences in the population, where the comparison is preferably performed by hybridising the labeled NAS for each of the
                                                                                                           each gene specific primer has a sequence complementary to a distinct mRNA, and each labeled NA is generated using a single gene specific primer. The method is useful for producing a sub-population of labeled
                                                                                                                                                                       The invention relates to producing a sub-population of labeled nucleic acids (NAs) comprising contacting a NA sample from a physiological source, with a pool of 50 distinct gene specific primers under suitable conditions to enzymatically generate sub-population of NAs, where
                                                                                                                                                                                                                                                                                                Producing sub-population of labeled nucleic acids, useful for analysing differences in RNA profiles between several different physiological sources, using set of distinct gene specific primers -
                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-314699/35
                                                                                                                                                                                                                                                                      Example 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Primer; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABK66892 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              bcr-abl, c-fgr and c-yes. The second oligonucleotide is specific for a nuclear oncogene or proto-oncogene selected from myc, jun, c-ets, c-fos, c-myb, B-myb, c-rel, c-vav, c-ski, c-spi, cyclin Dl, PML/RAR alpha, AMLI/MTGB, E2A/pri and ALL-1/AF-4. The composition is used for treating cancer. The combination of antisense oligonucleotides has synergistically enhanced ability to inhibit growth of cancer cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                               (CLON-) CLONTECH LAB INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US6352829-B1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABK66892;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 26 BP; 5 A; 4 C; 14 G; 3 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            specific for a cytoplasmic oncogene or proto-oncogene selected from ras, raf, EGF-1, c-fms, c-ros, c-kit, c-met, c-trk, c-src, c-abl,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           present invention. The present invention describes a composition which comprises two antisense oligonucleotides. The first oligonucleotide is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence represents an antisense oligonucleotide from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 4; Column 33; 92pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5 ATGTGACCGCCTCCCGCTCCCTCACC 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATGTGACCGCCTCCCGCTCCCTCACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    l Similarity
26; Conser
                                                                                                                                                                                                                                                                SEQ ID No 980; 11pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              specific PCR primer
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                                                                                                                                                                                                                                                                                                                                                                                                              Jokhadze
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            microarray; differential expression analysis; human
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Best Local :
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                        proteins, cytochromes, kinesins, cytokines, interferons, interleukins, G-protein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encode
                                                                                                                                                                                                                                                                                                                          Polymorphic nucleic acids encoding e.g. amylases, oncogenes and histones, useful for diagnosing and cancer, autoimmune diseases and infections -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1963 TCTCTACCGAAGATCAACCGGAGCGC 1988
                                                                                                                The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases, amyloid proteins, angiopoiet apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     interleukin; G-protein coupled receptor; thioesterase; multifactorial disease; autoimmune disease; infection; nervous system disease; ss.
                                                                                                                                                                                                                                                                  Claim 1; Page 2710; 4143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-DEC-1999; 99US-0173419
27-DEC-2000; 2000US-0173419
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-JAN-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            complement related protein; cytochrome; kinesin; cytokine; interferon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAL31396
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (CURA-)
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                                                                                                                                                                                                                                                                                                                                                         cyclins, polymerases, treating, e.g.
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s encoded
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                                                                                                                   The present invention describes human latexin protein 46 (I). (I) has cytostatic, antiviral, anti-HIV, immunomodulatory and antiinflammatory activities. (I) and the polynucleotide encoding it are applicable in the diagnosis and treatment of malignant tumour, haemopathy, HIV
                                                                                                                                                                                                                                                                                                     Human latexin protein 46 and encoded polynucleotide, applicable in diagnosis and treatment of malignant tumor, hemopathy, HIV infection, immunological diseases and inflammation -
                                                                                                                                                                                                                                                                                                                                                                                                           WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; latexin protein 46; cytostatic; antiviral; anti-HIV; immunomodulatory; antiinflammatory; diagnosis; malignant tu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                infection, immunological diseases and various inflammations. sequence represents a PCR primer for human latexin protein 4
                                                                                                                                                                                                                                                      Example 3; Page 18; 39pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DOOR GENE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             G; 28 T; 0
                                                 present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TECHNOLOGY LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    diagnosis; malignant tumour;
gical disease; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  , N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .3e+03;
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                                                                      ammations. The present protein 46, which is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  diabetes,
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C-raf;
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KW C-raf;
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KW G-raf;
XX GENE ET MISC_F
FT MIS
                                                                                     c-raf gene. The HPH molecules function through RNA/DNA inhibition. HPH molecule binds to and blocks the function of a target nucleic a and modulated cellular and viral processes such as splicing, editing translation. The HPH molecule can be used therapeutically, in target validation, to identify gene function and/or therapeutic targets, analysis of mutations in diseased cell and to detect specific RNA.
  greater
               hairpin structure improves resistance to nuclease degradation localization within the cell, and uptake by cells. The HPH may sequence that activates RNase H (for cleaving RNA) and its spe
                                                                                                                                                                                                                            molecule is used
c-raf gene. The H
                                                                                                                                                                                                                                         A method for modulating the function of a target sequence in a comprises treatment with a hairpin hybridizer (HPH) nucleic acid such as that represented by the present sequence. The present HP molecule is used in an example of the invention and targets the
                                                                                                                                                                                                                                                                                                                                                                           Claim 46; Page 76; 128pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Modulating target sequence in a ce
for identifying gene function, by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             misc_RNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (RIBO-)
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c-raf; inhibitor; detect; nuclease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hairpin hybridizer molecule #1 targeting c-raf RNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 cell, useful e.g. therapeutically by treatment with novel hairpin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 'n
sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hairpin hybridizer molecule; HPH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Jarvis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                internucleotide linkage
                                                                                                                                                                                                                                                                                             sequence in a cell
PH) nucleic acid mo
                  The HPH may include a and its specificity is
                                                                                                               t nucleic acid, ing, editing, ar y, in target targets, for
                                                                                                                                                                                                                                                                           HPH
                                                                                                                                                                                                                                                 human
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      c reaction into a double-stranded molecule. Primers #3 and #4, nested relative to primers #1 and #2 respectively, are used as tagging primers in the tagging reaction. A forward tagging primer has a defined sequence at its 5' end (+TAG sequence) while a reverse tagging primer has a different defined sequence at its 3' end (-TAG sequence). The purpose of the tagging reaction is to introduce the condition of the sequence to be amplified. The competitive amplification reaction involves using a single pair of generic primers, whose sequences are complementary to the +TAG and -TAG sequences, to amplify the different tagging step. This amplification reaction is competitive due to the use of a single primer pair to amplify the different target RNAS. Probe #5, complementary to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SS X
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Best Local
                                                                                                                                                                                                       the region of the resulting cDNA to be amplified during the MC reaction into a double-stranded molecule. Primers \#3 and \#4,
                                                                                                                                                                                                                                       by a tagging reaction and a competitive amplification reaction. RT-PCR reaction uses a primer #2 to convert target mRNA into cDJ Primer #1 in combination with primer #2 is then used to convert
                                                                                                                                                                                                                                                                                                                                        The present invention
                                                                                                                                                                                                                                                                                                                                                                                                                   Determining relative amounts of different nucleic acids by multiple: competitive polymerase chain reaction - involves tagging target and control sequences then amplification with generic primer pair corresponding to tagging sequences, used e.g. to determine response
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RT-PCR; tagginy ....
probe; housekeeping gene;
                                                                                                                                                                                                                                                                                                        by using the multiplex competitive PCR reaction (MC-PCR).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1998-447252/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Multiplex competitive PCR RT-PCR; tagging reaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C-raf probe
                                                                                                                                                                                                                                                                                     reaction involves a reverse-transcriptase (RT-PCR) reaction followed
                                                                                                                                                                                                                                                                                                                          relative amounts of two
                                                                                                                                                                                                                                                                                                                                                                        Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Thompson
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26; Conservative
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                                                                                                                                                                                                                                                                                                                                        provides a method for determining
                                                                                                                                                                                                                                                                                                                       or more different nucleic acid molecules
                                                                                                                                                                                                                                                                                                                                                                      English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       competitive amplification reaction: C-raf; hybridisation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        reaction; MC-PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ВP
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Pred. No. 3.7e+03
0; Mismatches
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                                                                                                                                                                                                                                                       into cDNA.
                                                                                                                                                                                                                         the MC-PCR
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  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   target mRNAs in a sample using the primer set #1-#5 for each target mRNA. In the example given, primers #1, #2, #3, #4 and probe #5 are the C-raf primers 1 (ANV33092), 2 (AAV33093), 3 (AAV33094), 4 (AAV33095) and probe 5 respectively. These primers/probes were used to illustrate the method of the invention. The method claims to allow detection of low-abundance mRNA in small samples (e.g. 10 ng is sufficient) with high precision, and uses housekeeping genes as controls for RNA input and integrity. Also, a large number of samples may be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA-RNA hybrid; phosphorothioate; hairpin hybridizer molecule; HPH;
c-raf; inhibitor; detect; nuclease resistance; gene function;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               misc_RNA
                                                                                                                                                                                                                                                                                                               misc_feature
                                                                                                                                                                                                                                                                                                                                                     misc_feature
                                                                                                                                                                                                                                                                                                                                                                                            misc_RNA
                                                                                                                                                                                                                                                                                                                                                                                                                      stem_loop
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene expression modulator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA30000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA30000 standard; DNA; 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 24 BP; 7 A; 9 C; 3 G; 5 T; 0 other;
Modulating target sequence in a cell, useful e.g. therapeutically or
                           WPI; 2000-292841/25
                                                    Hartmann T,
                                                                                                      21-SEP-1998;
                                                                                                                               20-SEP-1999;
                                                                                                                                                        30-MAR-2000.
                                                                                                                                                                                  WO200017346-A2
                                                                                                                                                                                                                                    misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hairpin hybridizer molecule #5 targeting c-raf RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            throughput screening, and does not require continuous monitoring.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the amplified product.
                                                                           (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             816 GCACAGATATTCTACACCTCACGC 839
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 GCACAGATATTCTACACCTCACGC 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                   Zwick M,
                                                                                                      98US-0101174
                                                                                                                               99WO-US21865
                                                                                                                                                                                                                                                                                                                /note=
12..20
                                                                                                                                                                                                                                                                                                                                       /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                    /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                               /note-
                                                                                                                                                                                                                                                note-
                                                                                                                                                                                                                                                                                     /note-
                                                                                                                                                                                                                                                                                                                                                                                /*tag=
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                                                                                                                                                                                                                                                                                                    *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%;
                                                  Thompson
                                                                                                                                                                                                                                                                                                                                                               "2'-0-methyl-ribonucleotides"
                                                                                                                                                                                                                                               "2'-O-methyl-ribonucleotides
                                                                                                                                                                                                                                                                                     "Phosphorothioate internucleotide linkage'
                                                                                                                                                                                                                                                                                                                           "Phosphorothioate internucleotide linkage"
                                                                                                                                                                                                         "Phosphorothioate internucleotide linkage"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The MC-PCR reaction can amplify one or more
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 24;
Pred. No.
                                                  ď,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                   Jarvis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 19;
3.3e+03;
                                                    7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene function;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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for identifying gene function, by treatment with novel hairpin hybridizer nucleic acid molecules
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such as that represented by the present sequence. The present HPH molecule is used in an example of the invention and targets the human c-raf gene. The HPH molecules function through RNA/DNA inhibition. The HPH molecule binds to and blocks the function of a target nucleic acid, and modulated cellular and viral processes such as splicing, editing, a translation. The HPH molecule can be used therapeutically, in target validation, to identify gene function and/or therapeutic targets, for analysis of mutations in diseased cell and to detect specific RNA. The hairpin structure improves resistance to nuclease degradation, localization within the cell, and uptake by cells. The HPH may include A method for modulating the function of a target sequence in a comprises treatment with a hairpin hybridizer (HPH) nucleic ac sequence that activates ${\tt RNASE}$ H (for cleaving ${\tt RNA}$) and its specificity greater than for linear antisense sequences. quence in a cell nucleic acid molecule нрн may include editing, and ĺS

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Sequence 31 BP; 6 A; 7 C; 11 G; 1 T; 6 U; 0 other;
 0.8%;
DB 21;
Length 31,
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Query Match Best Local Matches 2029 ATCAATGCTTGCACGCTGACCACG 2052 Similarity Conservative 100.0%; 0; Score 24; Pred. No. Mismatches 3.9e+03; 0 Indels 0; Gaps 0

. . . Db

27 ATCAATGCTTGCACGCTGACCACG 4

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RESULT 13
AAA29998/c
                                                                                                 AAA29998 standard; DNA; 33 BP
                                 09-AUG-2000
Hairpin
                                                                AAA29998
hybridizer molecule #3 targeting c-raf RNA.
                               (first entry)
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DNA-RNA hybrid; phosphorothicate; hairpin hybridizer molecule; c-raf; inhibitor; detect; nuclease resistance; gene function; gene expression modulator; ss. HPH;

				•
f Phosphorothioate internucleotide linkage"	f "Phosphorothioate in	/*tag= /note=		
		3033	misc_feature	7
leotides"	"2'-0-methyl-ribonucleotides"	/note=		7
	Ф	/*tag=		7
		2233	misc_RNA	~
"Phosphorothioate internucleotide linkage"	"Phosphorothioate in	/note=		-,
	٥	/*tag=		7
	•	1321	misc_feature	7
"Phosphorothioate internucleotide linkage"	"Phosphorothioate in	/note=		7
	C	/*tag=		7
		1 4	misc_feature	7
leotides"	<pre>/note= "2'-0-methyl-ribonucleotides"</pre>	/note=		7
	. 0	/*tag=		7
•		112	misc_RNA	_
	מ	/*tag=		_
		133	stem_loop	_
	Location/Qualifiers	Locatio	Key	-
			synthetic.	~ 0

21-SEP-1998; 20-SEP-1999; 30-MAR-2000 WO200017346-A2

98US-0101174 99WO-US21865

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AAA26239/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A method for modulating the function of a target sequence in a cell comprises treatment with a hairpin hybridizer (HPH) nucleic acid molecule such as that represented by the present sequence. The fresent HPH molecule is used in an example of the invention and targets the human c-raf gene. The HPH molecules function through RNA/DNA inhibition. The HPH molecule binds to and blocks the function of a target nucleic acid, and modulated cellular and viral processes such as splicing, editing, and translation. The HPH molecule can be used therapeutically, in target validation, to identify gene function and/or therapeutic targets. For
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         rranslation. The HPH molecule can be used therapeutically, in target validation, to identify gene function and/or therapeutic targets, for analysis of mutations in diseased cell and to detect specific RNA. The hairpin structure improves resistance to nuclease degradation, localization within the cell, and uptake by cells. The HPH may include a sequence that activates RNase H (for cleaving RNA) and its specificity is greater than for linear antiscance.
                                                      Thompson JD, Reynolds M,
                                                                                                                                                                                                                                                                                                     Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; hammerhead ribozyme; hairpin ribozyme; antisense ogene expression modification; cancer; phosphorothic
                                                                                                                                                                                                                                                                                                                                                           C-raf antisense oligonucleotide sequence SEQ
                                        Matulic-Adamic
                                                                                              (RIBO-) RIBOZYME
                                                                                                                           20-APR-1998;
23-JUN-1998;
                                                                                                                                                                       19-APR-1999;
                                                                                                                                                                                                   28-OCT-1999
                                                                                                                                                                                                                                W09954459-A2
                                                                                                                                                                                                                                                            Homo
                                                                                                                                                                                                                                                                                         anticancer; breast cancer; endometrium
                                                                                                                                                                                                                                                                                                                                                                                           19-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA26239 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2029 ATCAATGCTTGCACGCTGACCACG 2052
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 44; Page 76; 128pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Modulating target sequence in a c
for identifying gene function, by
hybridizer nucleic acid molecules
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (RIBO-) RIBOZYME PHARM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28
                                                                                                                                                                                                                                                           sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                   Beigelman L,
Zwick M, Jary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 6 A; 8 C; 12 G; 1 T; 6 U; 0 othe
                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zwick M,
                                                                                                                           98US-0082404
98US-0103636
                                                                                                                                                                     99WO-US08547
                                                                                              PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear antisense sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INC
                                                      Jarvis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Thompson
                                                                                                                                                                                                                                                                                                                                                                                                                                                  23
                                                   McSwiggen JA,
is T, Woolf T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                  ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 24; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cell, useful e.g. therapeutically or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               treatment with novel hairpin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Jarvis
                                                                                                                                                                                                                                                                                                     phosphorothicate;
                                                                                                                                                                                                                                                                                                                 antisense oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4.1e+03;
                                                   Karpeisky A,
Haeberli P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21;
                                                                                                                                                                                                                                                                                                                                                              ID NO:2737
                                                                                                                                                                                                                                                                                         SS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ength 33;
                                                                                                                                                                                                                                                                                                                                 cleavage;
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                                                                  Bellon
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Matches
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AAA26242/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                              Thompson JD, B
Reynolds M, Zw
Matulic-Adamic
                                                                                                                                                                                     20-APR-1998;
23-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            link, having endonuclease activity. (A), and more generally any catalytic nucleic acid (A') that modulates expression of the oestrogen receptor gene, are used to treat cancer (particularly of breast or endometrium), in vivo or by transforming cells ex vivo and implanting treated cells, or for other conditions associated with levels of cestrogen receptor. Because of the high selectivity for targeted RNA, (A)
                                                                                                                                                                                                                                                                                                                                                                                           Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavaye; hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide; gene expression modification; cancer; phosphorothioate; endonuclease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C-raf antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2029 ATCAATGCTTGCACGCTGACCAC 2051
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      restriction endonucleases are used with DNA). The combination of modifications in (A) improves resistance to nucleases, binding affinity and/or activity. AAA23503 to AAA24747 represent oestrogen receptor hammerhead ribozyme sequences, and AAA24748 to AAA25992 represent their corresponding target sequences. AAA2593 to AAA2505 represent oestrogen receptor hairpin ribozyme sequences, and AAA26107 to AAA26218 represent their corresponding target sequences. AAA26219 to AAA26271 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            can also be used to correlate inhibition of gene expression with alterations in phenotype, particularly for identification of therapeutic targets, and as research reagents (for RNA, in the same way that
                                                                                                                                                   (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                              anticancer; breast cancer; endometrium cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA26242 standard; DNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            other ribozyme sequences and antisense oligonucleotides used in the exemplification of the present invention.
                                                                                                                                                                                                                                       19-APR-1999;
                                                                                                                                                                                                                                                                         28-OCT-1999
                                                                                                                                                                                                                                                                                                           WO9954459-A2
                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA26242;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 23 BP; 5 A; 4 C; 8 G; 3 T; 3 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes nucleic acids (A) that interact
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New nucleic acids that interact, and optionally cleave, target sequences, used to treat cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
nes 23; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATCAATGCTTGCACGCTGACCAC 1
                                                                                 Zwick M,
nic J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                  Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence and contain at
                                                                                                                                                                                   98US-0082404
98US-0103636
                                                                                                                                                                                                                                       99WO-US08547.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                oligonucleotide sequence SEQ ID NO:2740.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     148pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.8%;
                                                                                                n L, McSwiggen JA,
Jarvis T, Woolf T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 23; pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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                                                                                                Karpeisky
Haeberli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   phosphoro
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                                                                                                 PΑ
                                                                                                                Bellon
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0;

New nucleic acids that interact, sequences, used to treat cancer

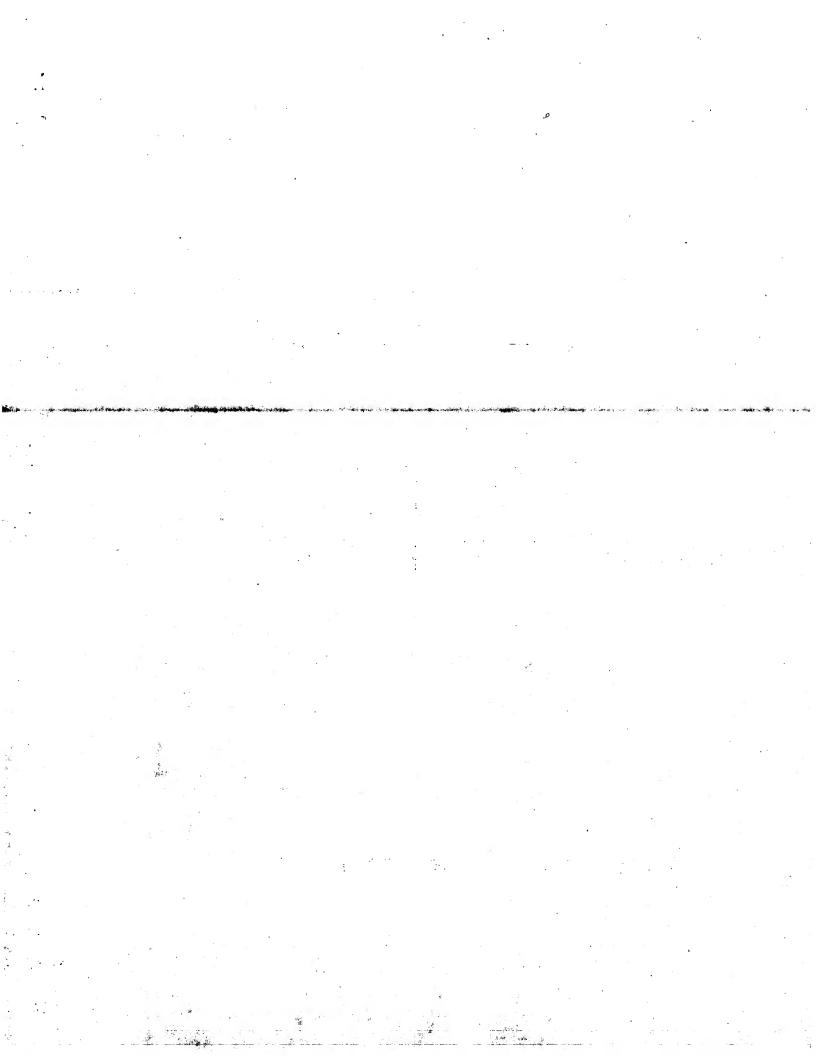
and optionally cleave,

target

WPI; 2000-013248/01.

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Search completed: May 16, 2003, 07:08:58 Job time : 644 secs
                                                                                                                                                                                                                                                                                    밁
                                                                                                                                     ş
                                                                                                                                                                                                                                                                                                              The present invention describes nucleic acids (A) that interact stably CC with a target sequence and contain at least one phosphoro(di)thioate CC link, having endonuclease activity. (A), and more generally any CC catalytic nucleic acid (A') that modulates expression of the oestrogen CC receptor gene, are used to treat cancer (particularly of breast or CC endometrium), in vivo or by transforming cells ex vivo and implanting CC treated cells, or for other conditions associated with levels of CC oestrogen receptor. Because of the high selectivity for targeted RNA, (A) CC can also be used to correlate inhibition of gene expression with CC alterations in phenotype, particularly for identification of therapeutic targets, and as research reagents (for RNA, in the same way that CC restriction endonucleases are used with DNA). The combination of CC modifications in (A) improves resistance to nucleases, binding affinity CA and/Or activity. ANA25503 to ANA24747 represent oestrogen receptor activity. ANA25503 to ANA24748 to ANA25992 represent their CC corresponding target sequences, and ANA26107 to ANA26271 represent CC their corresponding target sequences. ANA26219 to ANA26218 represent CC exemplification of the present invention.
                                                                                                                                                                              Query Match 0.8%; Score 23; DB Best Local Similarity 100.0%; Pred. No. 6.6 Matches 23; Conservative 0; Mismatches
                                                                                       Sequence 23 BP; 4 A; 8 C; 5 G; 2 T; 4 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 8; Page 105; 148pp; English.
                                                                                                                                                                                                      DB 21; I
. 6.6e+03;
                                                                                                                                                                                     0
                                                                                                                                                                                                                              Length 23;
                                                                                                                                                                                Indels
                                                                                                                                                                       . 0
                                                                                                                                                                                Gaps
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GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                              Result
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Title:
Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Minimum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Scoring table:
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    nucleic search, using sw model

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2977
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Gapop 10.0 , Gapext 1.0
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1: /cgn2_6/ptodata/
2: /cgn2_6/ptodata/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/BCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/backfiles1.seq:*
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                               US-09-482-084-2
US-08-850-049-17
US-08-850-049-17
US-07-858-7478-17
US-07-858-7478-17
US-09-614-117-17
US-09-614-825-9
US-08-874-825-25
US-08-63-824-9
US-08-63-824-25
US-08-63-824-25
US-09-231-303-9
US-08-874-825-26
US-08-874-825-26
US-08-874-825-26
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US-08-859-998-980
US-09-225-928-980
US-08-957-327-2
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US-08-306-691B-5
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US-08-785-247-31
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US-08-859-998-979
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8612.988 Million cell updates/sec
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                                                                           ; LENGTH: 42 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: DNA
US-08-785-247-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 13,
Patent No. 6
                                  Matches
                                                      Query Match
                                                                                                                                                        TELEFAX: 212-381-0526 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                            SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
916 GTCCACATGGTCAGCACCACGCTGCCTGTGGACAGCAGGATG 957
                                         Local
                                                                                                                                                         TELEPHONE: 212-381-0526
                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                           CITY: New York
STATE: New York
COUNTRY: U.S.A.
                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                   ZIP: 10036
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                                  38;
                                           Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Application US/08785247
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Zhang, Yuhua
                                  Conservative
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 Sequence 14, App 	Sequence 13, App	Sequence 12, Appl	Sequence 10, Appl	Sequence 9, Appl.	Sequence 8, Appl:	Sequence 7, Appl:	Seguence 6, Appl:	Sequence 5, Appl:	Sequence 4, Appl:	Sequence 3, Appl:	Sequence 2, Appl.	Sequence 11, Appl	Sequence 41, Appl	Sequence 11, Appl	Sequence 6, Appli	Seguence 3, Appli	Sequence 26, App.

ALIGNMENTS

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TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: ASSAY FOR IDENTIFYING AGENTS WHICH ACT ON TITLE OF INVENTION: CERAMIDE-ACTIVATED PROTEIN KINASE, KINASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/785,247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          E: Cooper & Dunham LLP
1185 Avenue of the Americas
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                                                                                                                                                                                                                                                                                                                                        48582-A/JPW/CCA
Score 35.6; DB Pred. No. 0.099; 0; Mismatches
                                       DB 3;
  4; Indels
                                         Length 42;
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Gaps
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GTCCACATGGTCAGCGTCACGCTACCGGTGGACAGCAGGATG 42

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Best Local Similarity 89.7%;
Matches 35; Conservative
                                                                                                                                                                                                              Sequence 14,
                                                                                                                                                                                            Patent No.
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                                                         CORRESPONDENCE ADDRESS:
                                                                          NUMBER OF SEQUENCES:
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                                                                                     ITLE OF INVENTION:
ITLE OF INVENTION:
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MEDIUM TYPE: Floppy disk
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CURRENT APPLICATION DATA:
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ADDRESSEE: Cooper & Dunham LLP
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TOPOLOGY: lir
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REGISTRATION NUMBER: 28,678
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CLASSIFICATION: 514
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New York
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                       E: Cooper & Dunham LLP
1185 Avenue of the Americas
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                                                                                                                                 Liu, Jun
Zhang, Yuhua
                                                                                                                                                                Kolesnick, Richard N.
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SYSTEM: PC-DOS/MS-DOS
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                                                                                     SUPPRESSOR OF RAS,
                                                                                               ASSAY FOR IDENTIFYING AGENTS WHICH ACT ON THE CERAMIDE-ACTIVATED PROTEIN KINASE, KINASE
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                                                                                SAID AGENTS
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Best Local Similarity 85.78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO:
    TELEPHONE: 415-322-5070
                                                            FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Field, Bret E.
                                                                                                                                                                                   SOFTWARE: FastSEQ for Windows Version CURRENT APPLICATION DATA:
                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
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ATTORNEY/AGENT INFORMATION:
NAME: White, John P. 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        916 GTCCACATGGTCAGCACCACGCTGCCTGTGGACAGCAGGATG 957
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                                                                                                                                                                                                                                                                                                                                                                                                 ITLE OF INVENTION:
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                               REFERENCE/DOCKET NUMBER:
                                               REGISTRATION NUMBER:
                                                                                                                                         CLASSIFICATION:
                                                                                                                                                     APPLICATION NUMBER: US/08/859,998 FILING DATE: 21-MAY-1997
                                                                                                                                                                                                    OPERATING SYSTEM:
SOFTWARE: FastSE
                                                                                                                                                                                                                                  COMPUTER:
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VENTION: METHOD OF ASSAYING DIFFERENTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chenchik, Alex
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                                                                                                                                                                                                                             IBM Compatible
                                                                                                                                                                                                                                                                                                                                    Fish & Richardson, P.C.
00 Sand Hill Road, Suite 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                Diskette
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Pred. No. 1;
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Length 42; Indels

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Gaps

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NFORMATION FOR SEQ ID NO:

SEQUENCE CHARACTERISTICS

ENGTH:

28 base pairs

1704 CAGTTTCCAGTCGGATGTCTACTCCTAT 1731

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OTHER INFORMATION: oligonucleotide primer; SEQUENCE DESCRIPTION: SEQ ID NO: 979: US-09-225-928-979
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              Query Match
Best Local Similarity
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Best Local Similarity
                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 979 SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 CAGTTTCCAGTCGGATGTCTACTCCTAT 28
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jokhadze, George
Bibilashvilli, Robert
TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
                                                                                                                           MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                  TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Chenchik, Alex
                                                                                                                                                   LENGTH: 28 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/09/225,928 FILING DATE: 05-Jan-1999 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 37,620
                                                                                                                                                                                                                                                                                                                 NAME: Field, Bret
                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 08/859,998 FILING DATE: 21-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: Windows95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Menlo Park
                                                                                                                                         ropology:
                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                       ELEFAX: 415-854-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OF SEQUENCES:
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              Score 28;
Pred. No.
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Pred. No.
 Mismatches
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                DB 4;
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RESULT 6
US-08-306-691B-5/c
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US-08-306-691B-6
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                                                                                                                                                                                                                                                                                                                                              US-08-306-691B-5
                                                                                                                                     Sequence 6, Applic Patent No. 5734039
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Best Local (
                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: NO. 5734039e INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Seidel, Gonda, Lavorgna & Monaco, P.C.
STREET: Two Penn Center, Suite 1800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: WordPerfect CURRENT APPLICATION DATA:
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                                                               TITLE OF INVENTION:
                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: (215) 568-8383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 514
PRIOR APPLICATION DATA:
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                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Philadelphia
STATE: Pennanca
                                                                                                                                                                                                                           26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Monaco, Daniel A. REGISTRATION NUMBER: 30,4 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PS, OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
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                                                                                                                                                   Application US/08306691B
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                                                                                         Skorski, Tomasz
                                                                                                                                                                                                                                                                                      Conservative
                                                                                                          Calabretta, Bruno
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                                                                                                                                                                                                                                                                                                                                                                          single stranded
                                                          ANTISENSE OLIGONUCLEOTIDES TARGETING COOPERATING ONCOGENES
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75;
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US-08-859-998-980/
Sequence 980, Application US/08859998
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NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                       ENERAL INFORMATION:
                                                                     CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                            ATTORNEY/AGENT INFORMATION:
                                                                                                                            SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 26 Nucleotides
                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS
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MEDIUM TYPE: Diskette, 3.50 inch, 720 Kb
                                                                                                                                                                                                                                           STREET: 2200 Sand Hill Road, Suite 100 CITY: Menlo Park
STATE: CA
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                           PPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (215)
NAME: Field, Bret E. REGISTRATION NUMBER: 37,620
                                            FILING DATE:
                                                         APPLICATION NUMBER:
                                                                                                APPLICATION NUMBER: US/0
FILING DATE: 21-MAY-1997
                                                                                                                                                           OPERATING SYSTEM:
                                                                                                                                                                          COMPUTER:
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                                                                                                                                                                                                                              COUNTRY:
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ilarity 100.0%;
Conservative (
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                                                                                                               US/08/859,998
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                                                                                                                                                                                                                                                                                                                                                          Robert
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                                                                                                                                                                                                                                                                                                                                             ASSAYING DIFFERENTIAL
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RESULT 9
US-09-225-928-980/c
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Best Local Similarity
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                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 980:
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OTHER INFORMATION: oligonucleotide primer
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TELEPHONE: 415-322-5070
TELEFAX: 415-854-0875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26 TCTCTACCGAAGATCAACCGGAGCGC 1
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                                                                                                                                                                                                                                                        APPLICATION NUMBER: 08/859,998
FILING DATE: 21-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: WI
SOFTWARE: FastSEQ fo
CURRENT APPLICATION DATA:
OTHER INFORMATION: oligonucleotide primer SEQUENCE DESCRIPTION: SEQ ID NO: 980:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 2200 Sand Hill Road, Suite 100
                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
                                      FEATURE:
                                                       MOLECULE TYPE: DNA
                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Chenchik, Alex
                                                                                       LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/225,928
FILING DATE: 05-Jan-1999
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            : 26 base pairs
nucleic acid
                   OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Menlo Park
STATE: CA
                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 94025
                                                                                                                                                                                                    TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER:
                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: US
                                                                                                                                                                                     ELEFAX: 415-854-0875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Jokhadze, George
Bibilashvilli, R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IBM Compatible
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%; Pred. No. 75;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Robert
                                                                                                                                                                                                                                            09096/002001
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JS-09-482-084-2/c
                                                                      Sequence 2, Application US/09482084 Patent No. 6333314 GENERAL INFORMATION:
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                      NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1963 TCTCTACCGAAGATCAACCGGAGCGC 1988
                                                                                                                                                                                                                                                                                                     HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: ka
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
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                                                                                                                                                                                            121 GCTGCATCAATGGAGCACATACAGG 145
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ZIP: 22031
                                                                                                                                                                    25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: FILING DATE: 24-OC1 CLASSIFICATION: 51
                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Hendricks, Gle
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Fairfax
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                ELEPHONE:
  TITLE OF
                                                                                                                                                                                                                                  Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               P.O. Box 2509
                                                                                                                                                                                                                                                                                                                                                                                                                  (703) 591-4470
(703) 591-4428
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rahman, Aqulur
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PatentIn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                   unknown
                                                      Kasid, Usha
              Dritschilo, Anatoly
Rahman, Aqulur
                                      Gokhale, Prafulla
                                                                                                                                                                                                                                                                                                                     DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-OCT-1997
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                                                                                                                                                                                                                                                   0.8%;
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Liposomes containng Oligonucelotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     32,535
                                                                                                                                                                                                                                   Score 25; ; Pred. No.
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Pred. No.
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US-08-850-049-17
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                                                                                                                                                                                                                                                                                       Patent No.
            COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY
                                                                                                                                                                                                                                                           APPLICANT:
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                                                                                                                                                 STREET:
                                                                                                           COUNTRY:
                                                                                                                                                                ADDRESSEE:
                                                                                              10154
                                                                                                           ÚSA
                                                                   FLOPPY DISK
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Sequence 17, Application US/08850049 Patent No. 5965726
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                                                                        CORRESPONDENCE ADDRESS:
                                                                                                          TITLE OF INVENTION:
                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                   121 GCTGCATCAATGGAGCACATACAGG 145
                                                                                                                                                                                                                                                                                                                                   25 GCTGCATCAATGGAGCACATACAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: US
ZIP: 22031
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: unknown MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 25 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Hendricks, Glenna REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/482,084 FILING DATE: 13-Jan-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: P.O. |
CITY: Fairfax
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 08/957,327 FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: kasid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Hendricks and Assoc.
                                    345 PARK AVENUE
                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; ---
                                                     & FINNEGAN
                                                                                                            INHIBITORY/INSTABILITY REGIONS OF mRNA
                                                                                                                             METHOD OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 591-4428
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                                                                                                                                                                                                                                                                                                                                                                                                                          Score 25;
Pred. No.
                                                                                                                           ELIMINATING
                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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US/08/850,049

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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                   tent No. 5972596
ENERAL INFORMATION:
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   2207 TCACAGGGCCTTAACTTCATGTTGCCTTCTTTTCTATC 2244
                                                                                   CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 43 BASE PAIRS
                                                                                                                                              CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
TITLE OF INVENTION:
UMBER OF SEQUENCES:
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PRIOR APPLICATION DATA:
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                                                                                                                  APPLICATION NUMBER: FILING DATE: 26-OC
                                                                                                                                                                               COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS
                                                                                                                                                                                                                                                                                                                ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: MORRY, MARY J.
REGISTRATION NUMBER: 34,398
REFERENCE/DOCKET NUMBER: 20
                                                                       PPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 TCTCAGATACCTAGCTTCATATTGCCTTATTGTCTACC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    17, Application US/08050478 o. 5972596
                                                                                                                                                                                                                                    10154 .
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                                                                                                                                                                                                                                                                                  NEW YORK
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                                                                                                                                                                                                                                                                    NEW YORK
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                                                                                                                                                               WORDPERFECT 5.1
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                                                                                                                                                                                                                                                                                                              MORGAN & FINNEGAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (212)758-4800
                                                          29-MAR-1993
                                                                                                                  26-OCT-1994
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27-MAR-1992
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                                                                                                                                                                             PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                      METHOD OF ELIMINATING
                                                                       PCT/US93/02908
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Pred. No.
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Best Local Similarity
Matches 28; Conser
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GENERAL INFORMATION:
                                                                                                                                                                                              INFORMATION FOR
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                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                  HYPOTHETICAL:
                                                                                                  MOLECULE TYPE:
                                                                                                                                                                                                                          TELEPHONE: (212)758-4800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: FLOPPY
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                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA
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CORRESPONDENCE ADDRESS:
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LENGTH: 43 BASE PAIRS
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                                                                                                                            STRANDEDNESS:
                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS
DENTIFICATION METHOD:
THER INFORMATION: OLIGONUCLEOTIDE FOR MUTATING
                                                      VAME/KEY:
                                                                                                                 COPOLOGY:
                                                                                                                                                             ENGTH:
                                                                                                                                                                                                                                                                        REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: UFILING DATE: 19920327
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                                    OCATION:
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                                                                                                                                             NUCLEIC ACID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
                                                                                                                                                                                                         (212)751-6849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PAVLAKIS, GEORGE N.; FELBER, BARBARA
                                                                                                                 UNKNOWN
                                                                                                                                                                                                                                                                                                                                                                                                                                   WORDPERFECT 5.1
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                                                                                                                               SINGLE
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Pred. No.
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US-09-414-117-17
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                                                                                                                       ;
US-09-414-117-17
                                                          Query Match 0.7
Best Local Similarity 73.7
Matches 28; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.:
Best Local Similarity 73.:
Matches 28; Conservative
                                                                                                                                                                                                                   NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ENERAL INFORMATION: APPLICANT:
2207 TCACAGGGCCTTAACTTCATGTTGCCTTCTTTTCTATC 2244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: WORDPERFECT 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/858,747
FILING DATE: 27-MAR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/02908
FILING DATE: 29-MAR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION: TELEPHONE: (212)758-4800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                            NTTORNEY/AGENT INFORMATION:
NAME: MORRY J.
REGISTRATION NUMBER: 34,398
REFERENCE/DOCKET NUMBER: 20:
                                                                                                                                                     TYPE: NUCLEIC ACID
STRANDEDNESS: SINGLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 345 PARK AVENUE
CITY: NEW YORK
STATE: NEW YORK
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PPLICATION NUMBER:
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                                                                                                                                                                                  43 BASE PAIRS
                                                                                                                                                                                                                                 (212)751-6849
                                                                                                                                        LINEAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MORGAN & FINNEGAN
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                                                                         Score 22; DB 4;
Pred. No. 1.9e+03;
                                                            Mismatches
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Post-processing: Minimum Match
Maximum Match
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Perfect score:
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seq length: 50
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/cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
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/cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*
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/cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
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   GenCore version 5:1.4_p5_4578 (c) 1993 - 2003 Compugen Ltd.
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10213.154 Million cell updates/sec
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 0 US-09-801-274-835
US-09-930-283A-2
US-09-9343-722-13
US-09-765-081-213
US-09-764-891-10177
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0 US-09-827-289-19
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0 US-09-827-289-23
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US-09-827-699-23
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US-10-103-906-4
US-10-103-906-4
US-10-078-949-3
US-10-078-949-3
US-10-078-949-5
US-10-078-949-7
US-09-9835-371-37
US-09-996-263-8
US-09-996-263-10
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                                                                                                  Sequence 835, Appli
Sequence 2, Appli
Sequence 17, Appl
Sequence 213, Appl
Sequence 10177. A
Sequence 580, Appl
Sequence 99, Appl
Sequence 19, Appl
Sequence 23, Appl
Sequence 23, Appl
Sequence 62, Appl
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APPLICANT: Cargill, Michele
APPLICANT: Lieland, James S.

APPLICANT: Lander, Eric S.

TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POI
FILE REFERENCE: 2825.2009-001
CURRENT APPLICATION UNMBER: US/09/801,274
CURRENT FILING DATE: 2001-03-07
PRIOR APPLICATION NUMBER: US 60/187,510
PRIOR FILING DATE: 2000-03-07
PRIOR FILING DATE: 2000-05-02
PRIOR APPLICATION NUMBER: US 60/206,129
PRIOR FILING DATE: 2000-05-22
NUMBER OF SEQ ID NOS: 1802
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 835
LENGTH: 31
                                                                              RESULT 2
US-09-930-283A-2/c
                                                                                                                                                                                                                     ; TYPE: DNA; ORGANISM: Homo sapiens US-09-801-274-835
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US-09-801-274-835
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Best Local S
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Gokhale, Prafulla
Dritschilo, Anatoly
Rahman, Aqulur
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                                                                      US/09930283A
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US-09-996-263-13
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US-09-996-263-10
US-10-057-550-2
US-10-057-550-3
US-10-057-550-9
US-10-057-550-19
US-10-057-550-20
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Pred. No. 13;
1; Mismatches
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US-09-943-722-17
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Publication No.
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                                                                                                                                                                                                                                                                                                                        ENERAL INFORMATION:
                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
                                                                                                                                                                                                                                                                                       APPLICANT:
APPLICANT:
                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                   TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         121 GCTGCATCAATGGAGCACATACAGG 145
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                                 OPERATING SYSTEM:
                                                    COMPUTER:
                                                                                                                      COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ANTI-SENSE: Y
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 1999-07-15
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: unknown
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Hendricks and Assoc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Liposomes containg Oligonucelotides NUMBER OF SEQUENCES: 3
ARE: WORDPERFECT 5.1
APPLICATION DATA:
                                                                                                     10154
                                                                                                                                    NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/09/930,283A FILING DATE: 16-Aug-2001 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (703) 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Hendricks, Glenna
REGISTRATION NUMBER: 32,535
REFERENCE/DOCKET NUMBER: ka
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                      Application US/09943722 No. US20020192660A1
                                                                                                                                                                   345 PARK AVENUE
                                                                                                                   USA
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                                                IBM PC COMPATIBLE
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100.0%; Pi
                               PC-DOS/MS-DOS
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Pred. No.
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7.1e+02;
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US-09-765-081-213
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                                                                                            Matches
                                                                                                                           Query Match
                                                                                                                                                                                                            SEQ ID NO 213
LENGTH: 21
                                                                                                          Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No. US2002(
                                                                                                                                                                                                                                             NUMBER OF SEQ ID I
                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/765,081
CURRENT FILING DATE: 2001-01-18
PRIOR APPLICATION NUMBER: US 60/176,861
PRIOR FILING DATE: 2000-01-19
                                                                                                                                                                                                                                                                                                                                       APPLICANT: Ireland, James S.
APPLICANT: Lander, Eric S.
TITLE OF INVENTION: HUWAN SINGLE NUCLEOTIDE POLYMORPHISMS
FILE REFERENCE: 2825.2008-001
                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Cargill, Michele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (212)751-684
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                              TYPE: DNA
                                                                                                                                                                          ORGANISM: Homo sapiens
                                                       1237 GGAGATGTTGCAGTAAAGATC 1257
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LENGTH: 43 BASE PAIRS
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TYPE: N
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REGISTRATION NUMBER: 34,398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US U:
FILING DATE: 27-MAR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 TCTCAGATACCTAGCTTCATATTGCCTTATTGTCTACC 39
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                                                                                        Conservative
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(212)751-6849
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                                                                                                        Score 20.6;
Pred. No. 1
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Pred. No. 9.
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Length 43;

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Gaps.

DB 10;

21; 0;

Matches

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us-09-263-959-580/c
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                                                             US-09-263-959-580
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                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Prior application data removed - consult PALM or NUMBER OF SEQ ID NOS: 10231
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/764,891
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and
FILE REFERENCE: PC006
                                                                                                                                                    NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2868 ACAGGTTTTCTTGATGATTTGGGTTTTAATTTTGTTTTTATTG 2910
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                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
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les 29; Conserv
                                                                                          STRANDEDNESS:
                                                                                                                                                                                                          NAME: McMasters, David D. REGISTRATION NUMBER: 33,963 REFERENCE/DOCKET NUMBER: 92
                                                                                                                                                                                                                                                         TORNEY/AGENT
                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                             TOPOLOGY:
                                                                                                                                                                                                                                                                           CLASSIFICATION:
                                                                                                                                                                                                                                                                                         FILING DATE:
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              Local Similarity
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                                                                                                        nucleic acid
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                                                                                                                     31 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6300 Columbia Center, 701 Fifth Avenue
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Koop, Ben F.
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              0.7%;
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              Score 20.4;
Pred. No. 2.
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Pred. No. 2
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US-09-827-289-19
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                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver. SEQ ID NO 19
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Best Local Similarity
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                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/827,289
CURRENT FILING DATE: 2001-04-05
                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 2000-0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 2000-03-24
PRIOR APPLICATION NUMBER: PCT/US00/07938
PRIOR FILING DATE: 2000-03-24
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                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: U.S. 60/194843
                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Exte
                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Abarzua, Patricio
TITLE OF INVENTION: Process for Allele Discrimination Using
                                                                                                                                                                                               TYPE: DNA
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                                                                                                                                                              FEATURE:
                                                                                                                                                                           ORGANISM: Artificial Sequence
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2893 TTAATTTTGTTTTTTTTGCACCTGACAAAATACAGTTATCT 2933
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RRENT APPLICATION NUMBER: US/10/085,906
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                                                     Similarity
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FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                  2000-04-05
                                                                                                                         Description of Artificial Sequence: Pl primer for use in allele discrimination
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                                                   Score 20.2; DB 10;
Pred. No. 3.7e+04;
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Pred. No. 2
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RESULT 11
US-10-103-906-4/c
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US-09-822-698A-62
; Sequence 62, Apr
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                                                        Sequence 4, Application US/10103906
Patent No. US20020156268A1
                                              GENERAL
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Best Local Similarity
Matches 28; Conserv
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APPLICANT: Abarzua, Patricio
TITLE OF INVENTION: Process for Allele Discrimination Using Primer
TITLE OF INVENTION: Extension
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No. US20020146750A1
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                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 09/538,913
PRIOR FILING DATE: 2000-03-30
NUMBER OF SEQ ID NOS: 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT FILING DATE: 2001-03-30
                                                                                                                                                                                                                                                                                                                                                          EQ ID NO 62
LENGTH: 48
TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: MUCIN-1 SFILE REFERENCE: DYX-015.1 US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: nucleotide sequence coding
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                                          INFORMATION
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Krotz, Achim n.
McElroy, Bethany
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Pred. No. 3
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Pred. No. 3.7e+04;
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                                                                                                                           GENERAL INFORMATION:
                                                                                                                                               Sequence 5, Application US/10078949 Patent No. US20020165189A1
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LENGTH: 20
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SEQ ID NO 4
LENGTH: 20
CURRENT APPLICATION NUMBER: US/10/078,949
CURRENT FILING DATE: 2002-02-20
PRIOR APPLICATION NUMBER: 09/479,783
PRIOR FILING DATE: 2000-01-07
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                                                                     TITLE OF INVENTION: Oligoribonucleotides And Ribonucleases For Cleaving RNA FILE REFERENCE: ISIS5027
                                                                                                        APPLICANT: Crooke, Stanley T.
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PRIOR FILING DATE: 1997-06-06
PRIOR APPLICATION NUMBER: 08/659,440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Crooke, Stanley T.
TITLE OF INVENTION: Oligoribonucleotides And Ribonucleases For Cleaving RNA
FILE REFERENCE: ISIS5027
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 09/479,783
PRIOR FILING DATE: 2000-01-07
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CURRENT FILING DATE: 2002-02-20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Methods for Removing Dimethoxytrityl Groups From TITLE OF INVENTION: Oligonucleotides
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CURRENT FILING DATE: 2002-03-22
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Best Local Similarity
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; Sequence 37, Application US/09835371
; Publication No. US20020187473A1
; GENERAL INFORMATION:
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 7
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           APPLICANT: UHLMANN, Eugen
APPLICANT: WILL, David W
APPLICANT: WILL, David W
TITLE OF INVENTION: POLYAMIDE NUCLEIC ACID DERIVATIVES,
TITLE OF INVENTION: PROCESSES FOR PREPARING THEM
FILE REFERENCE: 02481.1743 SEQUENCE LISTING
CURRENT APPLICATION NUMBER: US/09/835,371
CURRENT FILING DATE: 2001-04-17
NUMBER OF SEQ ID NOS: 53
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PRIOR FILING DATE: 2000-01-07
PRIOR APPLICATION NUMBER: 08/870,608
PRIOR FILING DATE: 1997-06-06
PRIOR APPLICATION NUMBER: 08/659,440
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CURRENT FILING DATE: 2002-02-20
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ORGANISM: Artificial Sequence
FEATURE:
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FILING DATE: 1997-06-06
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PatentIn Ver. 2.1
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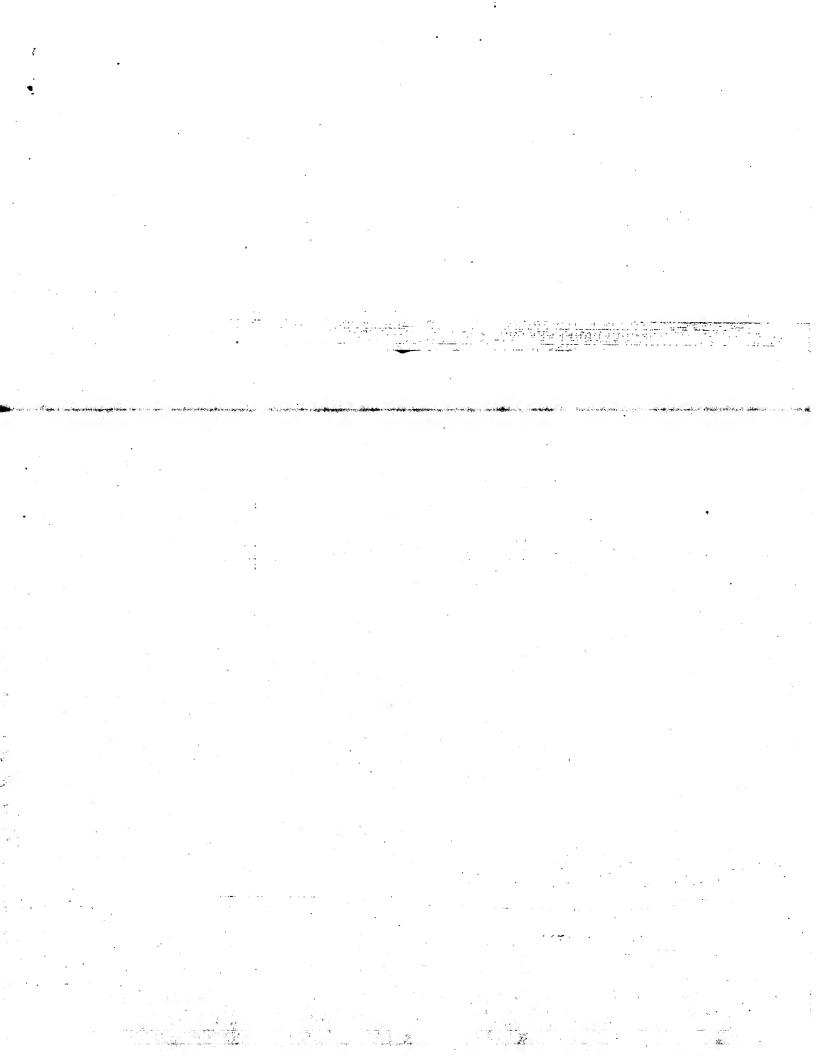
0;

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                                                                                                               ; OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: of PNA targeting CMV US-09-835-371-37
                         Q
                                                                                                                                                                                                    SEQ ID NO 37
LENGTH: 20
                                                                    Query Match
Best Local Similarity
                                                        Matches
                                                                                                                                                          TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                            2484 AATGCATGTCACAGGCGGGA 2503
20 AATGCATGTCACAGGCGGGA 1
                                                        Conservative
                                                                    100.0%;
                                                                   Score 20; Pred. No.
                                                        0
                                                                       No.
                                                                                DB 9; Length 20;
                                                       Indels
                                                                                                                                          base sequence
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Search completed: May 16, 2003, 12:48:11 : 381 secs

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Title:
Perfect score:
Sequence:
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Run on:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Minimum DB seq length: 0 Maximum DB seq length: 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OM nucleic - nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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2977
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /cgn2_6/ptodata/1/pna/US06_COMB.seq:*
                                       /cgn2_6/ptodata/1/pna/US099C_COMB.seq:
/cgn2_6/ptodata/1/pna/US099D_COMB.seq:
/cgn2_6/ptodata/1/pna/US100A_COMB.seq:
                   cgn2_6/ptodata/1/pna/US100B_COMB.seq:
cgn2_6/ptodata/1/pna/US101A_COMB.seq:
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                                                                                                                                                               //ptodata/1/pna/US098B_COMB.seq:*
//ptodata/1/pna/US098C_COMB.seq:*
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ptodata/1/pna/US085_COMB.seq:*
                                                                                                                /ptodata/1/pna/US099A_COMB.seq:
/ptodata/1/pna/US099B_COMB.seq:
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/ptodata/1/pna/US087_COMB.seq:*
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ptodata/1/pna/US096C_COMB.seq:
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ptodata/1/pna/US096A_COMB.seq:
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10_COMB.seq:*
11_COMB.seq:*
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/cgn2_6/ptodata/1/pna/US6041_COMB.seq:*
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/cgn2_6/ptodata/1/pna/US6040_COMB.seq:*
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                                                                                        cgn2_6/ptodata/1/pna/US6036_COMB.seq:
                                                     cgn2_6/ptodata/1/pna/US6038_COMB.seq:*
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4_COMB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMAR	
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	No.	Score	Match L	Length DB	œ	ID	Description
	c 1	50	1.7	50	1:	PCT-US01-47856-3119.	Sequence 3119, Ap
	c 2	50	1.7	50	40	US-10-131-827-3119 .	Sequence 3119, Ap
	ი 3	50	1.7	50	40	US-10-131-831-3119	Sequence 3119, Ap
	4	31.6	1.1	48	69	US-60-253-651-9462 :	Sequence 9462, Ap
	ر ت	30.6	1.0	31	31	US-09-801-274-835	Sequence 835, App
٠	6	30.6	1.0	-	69	US-60-253-653-25508 ;	Sequence 25508, A
	7	28	0.9	_	16	US-09-225-201B-979	Sequence 979, App
	ი 8	26		26	7	US-08-306-691-5	Sequence 5, Appli
		26	0.9	26	7	US-08-306-691-6	Sequence 6, Appli
	c 10	26		26	12	US-08-821-015-5	Sequence 5, Appli
	11	26	0.9	26	12	US-08-821-015-6	Seguence 6, Appli
	c 12	26		26	16	US-09-225-201B-980	Seguence 980, App
	c 13	25		25	17	US-09-354-109-2	Sequence 2, Appli
	14	25		25	17	US-09-396-196F-44994	Sequence 44994, A
	15	. 25		25	17	US-09-396-196F-44995 .	Sequence 44995, A
•	16	25	0.8	25	17	US-09-396-196F-44996	Sequence 44996, A
	17	25	0.8	25 .	17	US-09-396-196F-44997	Sequence 44997, A
	18	25	0.8	25	17	US-09-396-196F-44998	Sequence 44998, A
-	19	25	0.8	25	17	US-09-396-196F-44999	Sequence 44999, A
	20	25	0.8	25	17	US-09-396-196F-45000	Seguence 45000, A
	21	25	0.8	25	17	US-09-396-196F-45001	Sequence 45001, A

US-09-396-196F-45003
US-09-396-196F-45004
US-09-396-196F-45006
US-09-396-196F-45006
US-09-396-196F-45007
US-09-396-196F-45010
US-09-396-196F-45011
US-09-396-196F-45011
US-09-396-196F-45012
US-09-396-196F-45013
US-09-396-196F-45013
US-09-396-196F-45015
US-09-396-196F-45015
US-09-396-196F-45015
US-09-396-196F-44994
US-09-396-196G-44999

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TTGTTTTTATTGCACCTGACAAATACAGTTATCTGATGGTCCCTCAATT 1
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59148,
59150,
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APPLICANT: WOODWARD, ROBERT

APPLICANT: Ly, Ngoc

ITILE OF INVENTION: CHRONIC INFLAMMATORY DISEASES

FILE REFERENCE: 506612000120

CURRENT APPLICATION NUMBER: US/10/131,827

CURRENT FILING DATE: 2002-09-06

PRIOR APPLICATION NUMBER: US 10/006,290

PRIOR FILING DATE: 2001-10-22

PRIOR APPLICATION NUMBER: US 60/296,764

PRIOR FILING DATE: 2001-06-08

NUMBER OF SEQ ID NOS: 9090

SOFTWARE: PATEORITION OF SEQ ID NOS: 9090

SOFTWARE: DATE: 1005: 9090

SOFTWARE: DATE: 1007: CARPICAL OF SEQ ID NOS: 9090

TYPE: DNA

CDDANIEW. Homo Captions
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GENERAL INFORMATION:
APPLICANT: Glenn, Mattew
APPLICANT: Grigor, Murray R
TITLE OF INVENTION: Compositions isolated from bovine tissue
TITLE OF INVENTION: and methods for their use.
FILE REFERENCE: 1055P3
                                                                                                                                                                                         Sequence 9462, Application US/60253651
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 1.7
Best Local Similarity 100.
Matches 50; Conservative
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APPLICANT: Wohlgemuth, Jay
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR FILING DATE: 2001-06-08
NUMBER OF SEQ ID NOS: 9190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 10/006,290 PRIOR FILING DATE: 2001-10-22
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CURRENT APPLICATION NUMBER: US/10/131,831
CURRENT FILING DATE: 2002-08-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Ly, NGOC
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING AND MONITORING
TITLE OF INVENTION: TRANSPLANT REJECTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                        2899 TTGTTTTTATTGCACCTGACAAAATACAGTTATCTGATGGTCCCTCAATT 2948
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                                                                                                                                                                                                                                                                                                                                                         50 TIGTTTTTATTGCACCTGACAAAATACAGTTATCTGATGGTCCCTCAATT 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wohlgemuth, Jay
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fry, Kirk
Woodward, Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 50;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 40;
0.017;
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0.017;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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PCT-US01-47856-3119/c : Sequence 3119, Application PC/TUS0147856

ALIGNMENTS

APPLICANT: BIOCARDIA,

INC

Wohlgemuth, Jay Quertermous, Thomas Johnson, Frances

INFORMATION:

US-10-131-827-3119/c; Sequence 3119, Appli; GENERAL INFORMATION:

Application US/10131827

RESULT 2

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50

2899

; TYPE: DNA; Homo sapiens PCT-US01-47856-3119

LENGTH: 50

Best Local Similarity Matches 50; Conserv

Conservative

1.7%; Score 50; 100.0%; Pred. No. 0;

DB 1; 0.017;

Mismatches

0;

Query Match

CURRENT APPLICATION NUMBER: PCT/US01/47856
CURRENT FILING DATE: .2001-10-22
PRIOR APPLICATION NUMBER: US 60/241,994
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR FILING DATE: 2001-06-08
NUMBER OF SEQ ID NOS: 8832
SOFTWARE: Patentin version 3.1
SEQ ID NO 3119

APPLICANT: Matcuk, George
APPLICANT: Printips, Julie
APPLICANT: Woodward, Robert
APPLICANT: Woodward, Robert
APPLICANT: Ly, Ngoc
APPLICANT: Altman, Peter
TITLE OF INVENTION: LEUKOCYTE EXPRESSION PROFILING
FILE REFERENCE: 506612000140

APPLICANT: APPLICANT: APPLICANT: APPLICANT:

Fry,

CURRENT APPLICATION NUMBER: US/60/253,651 CURRENT FILING DATE: 2000-11-28 NUMBER OF SEQ ID NOS: 27858 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 9462

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; LENGTH: 48
; TYPE: DNA
; ORGANISM: Bovine
US-60-253-651-9462
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                                                                                                  NUMBER OF SEQ ID NOS: 30124
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 25508
LENGTH: 46
TYPE: DNA
                                                                                                                                                                                                                                  Sequence 25508, Application US/602
GENERAL INFORMATION:
APPLICANT: Glenn, Matthew
TITLE OF INVENTION: Compositions
TITLE OF INVENTION: and methods
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 835, Applica GENERAL INFORMATION:
                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 835
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
      Matches
                                                                     ORGANISM: Ovine
-60-253-653-25508
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -09-801-274-835
                                                                                                                                                                                                                                                                                                                       60-253-653-25508
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/206,129
PRIOR FILING DATE: 2000-05-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/801,274
CURRENT FILING DATE: 2001-03-07
PRIOR APPLICATION NUMBER: US 60/187,510
PRIOR FILING DATE: 2000-03-07
                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/60/253,653
CURRENT FILING DATE: 2000-11-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                      FILE REFERENCE: 1057P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Cargill, Michele APPLICANT: Ireland, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PPLICANT: Iréland, James S.
PPLICANT: Lander, Eric S.
ITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
ILE REFERENCE: 2825.2009-001
                                                                                                                                                                                                                                                                                                                                                                                                                        2144 CCCTTTCTCCAGAGGCAGAACACATGTTTTC 2174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1688 AGGATAACAACCCATTCAGTTTCCAGTCGGATGTCTACTCCTATGG 1733
                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
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     Conservative
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                    1.0%;
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                    Score 30.6;
Pred. No. 2.
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Pred. No. 2
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Pred. No. 1.5e+03
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                   .8e+03;
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                                    DB 69;
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                                  Length 46;
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OTHER INFORMATION: C;
SEQUENCE DESCRIPTION: SEQ
US-09-225-201B-979
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                                                                                                                                                                 US-08-306-691-5/c
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                                                                                                                                                                                        RESULT 8
                                                                                                                             Sequence 5, Application US/08306691 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                               1704 CAGTTTCCAGTCGGATGTCTACTCCTAT 1731
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 979:
                                                                                              APPLICANT:
                                                         CITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                            1 CAGTTTCCAGTCGGATGTCTACTCCTAT 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: US
ZIP: 94025
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/859,998 FILING DATE: 21-MAY-1997 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bibilashvilli, Robert
TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Chenchik,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CCACTCCCATTGGATCACGCTCCTTCGGGACTGTTTATAACGGCA
                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/225,201B FILING DATE: 05-Jan-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 37,620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 415-854-0875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Field,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Fish & Richardson, P.C. STREET: 2200 Sand Hill Road, Suite 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ENGTH: 28 base pairs
                                                                                        Skorski,
                                                                                                               Calabretta, Bruno
                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                              0.9%; 5c.
100:0%; Pr
                                                         ANTISENSE .
OLIGONUCLEOTIDES TARGETING COOPERATING ONCOGENES
Gonda, Lavorgna & Monaco, P.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bret E
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                                                                                                                                                                                                                                                                                                                                     Score 28; pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                              oligonucleotide primer Q ID NO: 979:
                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                      Length 28
                                                                                                                                                                                                                                                                                                                     Indels
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COMPUTER READABLE FORM:

: Two Penn Center, Suite 1800 Philadelphia

Pennsylvania

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us-08-306-691-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                           equence 6, Applicati
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX: None
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                  OPERATING SYSTEM: MS-DOS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
                         ATTORNEY/AGENT INFORMATION:
                                                                         CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 720 Kb
                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ELECOMMUNICATION INFORMATION: TELEPHONE: (215) 568-8383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                  ITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26
                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM:
SOFTWARE: WordPer
                                                           APPLICATION NUMBER:
                                                                                                                       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                      ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Monaco, Daniel A. REGISTRATION NUMBER: 30,480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ICATION NUMBER:
                                                                                                                                                                                                                               19102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                      Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                            Application US/08306691
                                                                                                                                                                                                                                                          Pennsylvania
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                                                                                                                                                                                                                                          U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (215) 568-5549
                                                                                                                                                                                                                                                                                                                                                                                            Calabretta, Bruno
                                                                                                                                                                                IBM PS/2
                                                                                                                                                                                                                                                                                                    Seidel,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E: Diskette, 3.50 inch, 720 Kb
IBM PS/2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    single stranded
                                                                                                                                                                                                                                                                                                                                                                  ANTISENSE
                                                                                                                                                                                                                                                                                                                                               OLIGONUCLEOTIDES TARGETING COOPERATING ONCOGENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.9%; Score 26;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                    Gonda,
                                                                                                                      US/08/306,691
A.
30,480
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                                                                                                                                                                                                                                                                                     , Lavorgna & Monaco, P.C. Suite 1800
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  Matches
                             Query Match
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                                                                                                                           INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: MS-DOS
SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                     TELECOMMUNICATION INFORMATION: TELEPHONE: (215) 568-8383
                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                         FILING DATE: September 15, 1994 ATTORNEY/AGENT INFORMATION:
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                                                                                      STRANDEDNESS:
                Local Similarity
                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PS/2
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX:
                                                                                                                 CENGTH:
                                                                                                                                                                       TELEFAX:
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                                                                                                                                                                                                                               REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE:
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                                                                                                  nucleic acid
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                                                                                                                                                                                                                                                Monaco,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pennsylvania
                                                                                                               26 Nucleotides
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Calabretta, Bruno
                                                                         linear
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                                                                                                                                                                                                                      Daniel A. 30,480
                                                                                  single stranded
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              0.9%;
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100.0%; Pred. No.
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                Pred. No.
 Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                       720 Kb
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3.6e+04
                             DB 12;
                3.6e+04
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                          Length 26:
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GENERAL INFORMATION:
APPLICANT: Chenchik, Alex
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: September 15, ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Diskette, 3.50 inch, 720 Kb
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                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish &
 COMPUTER READABLE FORM:
                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                        Bibilashvilli, Robert
TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
                                                                                                                                                                                                                                                                                       ATGTGACCGCCTCCCGCTCCCTCACC 26
                                                                                                                                                                                                                                                                                                                                                                                                                                        nucleic acid
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               COUNTRY: US
ZIP: 94025
                                                         CITY: Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          None
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                                                                                                                                                                                                                                                                                                                                                                                                                      linear
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                                                                                                                                                       Jokhadze, George
Bibilashvilli, Ro
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                                                                     E: Fish & Richardson, P.C. 2200 Sand Hill Road, Suite 100
                                                                                                                                                                                                                                                                                                                                                                                                                                  single stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Daniel A.
                                                                                                                                                                                                                                                                                                                                                             100.0%;
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                                                                                                                              EXPRESSION
                                                                                                                  1375
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8321-8 DI1
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Suite 1800
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ELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                  STREET: P.O. I
                                                                                                               CLASSIFICATION:
                                                                                                                                                                                    OPERATING SYSTEM:
                                                                                                                              FILING DATE:
                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                    COMPUTER:
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INFORMATION FOR SEQ ID NO: 980:
                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/957,327
FILING DATE: 24-OCT-1997
ATTORNEY/ACENT INFORMATION:
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                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
REGISTRATION NUMBER: 32,
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FILING DATE: 21-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Field, Bret E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
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SEQUENCE DESCRIPTION: SEQ
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Diskette
COMPUTER: IBM Compati
                                          Hendricks, Glenna
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REGISTRATION NUMBER: 37,620
PROPERIOR /PACKET NUMBER: 09096/002001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dritschilo,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gokhale, Prafulla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kasid, Usha
                                                                                                                                                                                                                                            IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                Hendricks and Assoc
                                                                                                                                                                                                                                                                                    Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                               Box 2509
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100.0%; Pred. No.
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Q ID NO: 980:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3.6e+04;
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Gaps

0

TELEFAX: (703) 591-4428 INFORMATION FOR SEQ ID NO: 2:

SEQUENCE CHARACTERISTICS:

nucleic acid

STRANDEDNESS:

unknown

single

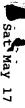
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RESULT 15
US-09-396-196F-44995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; SOFTWARE: FastSEQ for Windows Version 4.0; SEQ ID NO 44994; LENGTH: 25; TYPE: DNA; ORGANISM: mus musculus US-09-396-196F-44994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               밁
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US-09-354-109-2
                                                                                                                                                                                                                                                                      Sequence 44995, Application US/09396196F GENERAL INFORMATION:
                 PRIOR APPLICATION NUMBER: 60/100,678
PRIOR FILING DATE: 1998-09-17
NUMBER OF SEQ ID NOS: 127806
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4495
LENGTH: 25
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.8%;
Best Local Similarity 100.0%;
Matches 25; Conservative
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Best Local Similarity 100.
Matches 25; Conservative
                                                                                                                           FILE REFERENCE: 3101.1
CURRENT APPLICATION NUMBER: US/09/396,196F
CURRENT FILING DATE: 2001-09-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/396,196F CURRENT FILING DATE: 2001-09-15
                                                                                                                                                                           APPLICANT: David Mack
APPLICANT: David Lockhart
APPLICANT: Affymetrix, Inc.
AFFYMETRIX: Methods of Genetic Analysis
                                                                                                                                                                                                                                                    APPLICANT: Michael Mittmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/100,678 PRIOR FILING DATE: 1998-09-17
ORGANISM: mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 127806
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ITLE OF INVENTION: Methods of Genetic Analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: unl
MOLECULE TYPE:
HYPOTHETICAL:
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David Lockhart
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100.0%; Pr
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6.5e+04;
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US-09-396-196F-44995
Query Match 0.8%; Score 25; DB 17; Best Local Similarity 100.0%; Pred. No. 6.5e+04; Matches 25; Conservative 0; Mismatches 0;
                                     Length 25;
0;
Gaps
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B 1876 AGGCTGGTAGCTGACTGTGTGAAGA 1900 1 AGGCTGGTAGCTGACTGTGAAGA 25

Search completed: May 16, 2003, 12:16:27 Job time: 6348 secs



10 11 11 11 11 11 11 11 11 11 11 11 11 1	1 W 4 V 0 V 8 Q	Result No. 'S	Pred scor and	Database :	Post-proce	Minimum DB Maximum DB	Searched: Total numb	Scoring tab	Title: Perfect sc Sequence:	Run on:	OM nucleic	
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109044 714	Sequence 320641, Sequence 320641, Sequence 356623, Sequence 917222, Sequence 397923, Sequence 5803, Ap Sequence 1070, Ap	Description sequence 3119, Ap	ce to have a lt being printed, bution.	@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@			58442		aattaaattt 2977	Seconds ents) on cell updates/sec		
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0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7
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US-10-325-899-3367 US-10-325-899-4603	US-10-287-787-21970	US-10-325-899-1696 US-10-287-787-21968	US-09-912-293-43884 .	US-10-287-787-19197	US-10-325-899-7636	US-60-427-836-451365	US-60-427-808-943561	US-60-427-808-630023	US-60-427-808-506441	US-10-325-899-8045	US-10-325-899-3101	US-09-912-293-242736	US-10-349-143-2948	US-10-287-787-19790	US-10-287-787-19789	US-10-389-048-4355	US-10-287-787-4936	US-10-287-787-4937	US-09-912-293-130705 ·	US-60-427-808-116035
Sequence 3367, Ap Sequence 4603, Ap		Sequence 21968, A		Sequence 19197, A	Sequence 7636, Ap	Sequence 451365,	·Sequence 943561,	Sequence 630023,	Sequence 506441,	Sequence 8045, Ap	Sequence 3101, Ap	Sequence 242736,	Sequence 2948, Ap	Sequence 19790, A	Sequence 19789, A	Sequence 4355, Ap	Sequence 4936, Ap	Sequence 4937, Ap	Sequence 130705,	Sequence 116035,

ALIGNMENTS

US-10-325-899-3119/c
US-10-325-899-3119/c
; Sequence 3119, Application US/10325899
; GENERAL INFORMATION:

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PRIOR APPLICATION NUMBER: US 10/131,831
PRIOR FILING DATE: 2002-04-24
NUMBER OF SEQ ID NOS: 9966
SOFTWARE: Patentin version 3:1
SEQ ID NO 3119
LENGTH: 50
TYPE: DNA
ORGANISM: Homo sapiens
US-10-325-899-3119
                                                                                                                   US-10-075-994A-2/c ; Sequence 2, Application US/10075994A; GENERAL INFORMATION:
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Best Local Similarity 100.0%;
Matches 50; Conservative (
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APPLICANT: Wohldemuth, Jay
APPLICANT: Fry, Kirk
APPLICANT: Ly, Ngoc
APPLICANT: Woodward, Robert
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING AND MONITORING TRANSPL
TITLE OF INVENTION: REJECTION
FILE REFERENCE: 506612000122
CURRENT APPLICATION NUMBER: US/10/325,899
CURRENT FILING DATE: 2002-12-20
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR FILING DATE: 2001-06-08
PRIOR APPLICATION NUMBER: US 10/006,290
PRIOR FILING DATE: 2001-06-22
PRIOR FILING DATE: 2001-10-22
PRIOR FILING DATE: 2001-10-22
APPLICANT: KASID, Usha
APPLICANT: GOKHALE, Prafulla
APPLICANT: PIE, Jin
APPLICANT: MEMANI, Rajshree
APPLICANT: AHMAD, Imran
                                                                                                                                                                                                                                                      Score 50; DB 9; Ler
; Pred. No. 0.00061;
...amatches 0;
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; ORGANISM: Mus musculus 
US-60-427-808-556623
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                                            Query Match
Best Local
                                Matches
                                                                                                                                                    NUMBER OF SEQ ID NOS: 982914
SOFTWARE: Microarray Probe S
SEQ ID NO 556623
                                                                                                                                                                                                                                                           Sequence 556623, Ap
GENERAL INFORMATION
APPLICANT: Xue Mei
                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
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APPLICANT: Xue Mei Zhou
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SEQ ID NO 2
LENGTH: 25
                                                                                                                       LENGTH: 25
TYPE: DNA
                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/60/427,808
CURRENT FILING DATE: 2002-11-20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Methods of Genetic Analysis FILE REFERENCE: 3528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 982914
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/075,994A
CURRENT FILING DATE: 2002-02-15
NUMBER OF SEQ ID NOS: 7
2214 GCCTTAACTTCATGTTGCCTTCTTT 2238
                                                                                                                                                                                                                             ITLE OF INVENTION: Methods of Genetic Analysis of Mouse ILE REFERENCE: 3528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: RAHMAN, AGULIUT
TITLE OF INVENTION: CHEMOSENSITIZING WITH LIPOSOMES CONTAINING OLIGONUCLEOTIDES
FILE REFERENCE: 219604
                                                                                                                                                                                                                                                                                                                                                                          2210 CAGGGCCTTAACTTCATGTTGCCTT 2234
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                                Conservative
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                                            Pred. No.
                                                         Score 23.4;
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0; Mismatches
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Pred. No. 2
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Pred. No.
                           Mismatches
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                                           2.4e+04;
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                           Indels
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; ORGANISM: Rattus norvegicus
US-60-427-836-397923
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                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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SEQ ID NO 917222
LENGTH: 25
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Best Local Similarity
                                              FILE REFERENCE: 506612000122
CURRENT APPLICATION NUMBER: US/10/325,89:
CURRENT FILING DATE: 2002-12-20
                                                                                         APPLICANT: Ly, Ngoc
APPLICANT: Woodward, Robert
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING AND MONITORING TRANSPL
TITLE OF INVENTION: REJECTION
                                                                                                                                                                    APPLICANT: Wohlgemuth, Jay APPLICANT: Fry, Kirk
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                   PRIOR FILING DATE:
                                PRIOR APPLICATION NUMBER: US 60/296,764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/60/427,836 CURRENT FILING DATE: 2002-11-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Microarray Probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 699466
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Xue Mei Zhou
TITLE OF INVENTION: Methods of Genetic Analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILE REFERENCE: 3528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 98291
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APPLICATION NUMBER: US 10/006,290
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                   2001-06-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Methods of Genetic Analysis of
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Pred. No. 2
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PRIOR FILING DATE:

PRIOR APPLICATION NUMBER: US 10/131,831

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TCGCCCACCGAGATATTAAATCCAGCCACATAATGCTCTTTGA 45

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CURRENT APPLICATION NUMBER: US/60/
CURRENT FILING DATE: 2002-12-19
NUMBER OF SEQ ID NOS: 7137
SOPTWARE: PatentIn version 3.1
SEQ ID NO 1070
LENGTH: 43
TYPE: DNA
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Best Local Similarity
Matches 29; Conser
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 5803
                                                               Best Local Similarity
                                                                                                                                                     SEQ ID NO 19321
LENGTH: 45
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Best Local Similarity
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                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/60/288,292
CURRENT FILING DATE: 2001-05-02
                                                                                                                                                                                                                                                           TITLE OF INVENTION: Compounds
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                                                                                                                                                                                    NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
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                                                                                                                                                                                                                                                                                           APPLICANT: Glenn, Matthew APPLICANT: Norriss, Michael G
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Roemer, Terry
                                                                                                                                         TYPE: DNA
1520 TCATCCATAGAGACATGAAATCCAACAATATATTTCTCCCATGA 1562
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                                               30;
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Boone, Charles
                                               Conservative
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Pred. No.
                                                            Score 22.2; DB 11
Pred. No. 7.1e+04;
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                                              Mismatches
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                                              Indels
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PCT-US02-14877A-875

; Sequence 875, Application PC/TUS0214877A

; GENERAL INFORMATION:
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US-10-293-338-4068
                                                         RESULT 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; ORGANISM: Homo sapiens
US-10-293-338-4068
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SEQ ID NO 3876
LENGTH: 47
TYPE: DNA
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CURRENT FILING DATE: 2002-11-14
NUMBER OF SEQ ID NOS: 8785
SOFTWARE: PatentIn version 3.1
SEQ ID NO 4068
LENGTH: 22
TYPE: DNA
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GENERAL INFORMATION:
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PRIOR APPLICATION NUMBER: US
PRIOR FILING DATE: EARLIER FILING DATE: 1999-04-21
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US
PRIOR FILING DATE: EARLIER FILING DATE: 1998-11-23
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TITLE OF INVENTION: BIOINFORMATICALLY DETECTABLE GROUP OF NOVEL REGULATORY GENES
TITLE OF INVENTION: THEREOF
                                                                                                                                                                                                                                                                                                       FEATURE:
NAME/KEY: allele
LOCATION: 24
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CURRENT FILING DATE: 2003-01-21
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                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TLE OF INVENTION: Biallelic markers for use
                                                                                                                                                   2554 AACAGCCCAGGACTGCCGAGACTCTGGCCGCCGAAGGAGCCTGCT 2599
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                                                                                                                 AACAGCCTTGGGCTCACGTTACARCAGCATCCCCAAGGAGCCTCCT 46
                                                                                                                                                                                                             Similarity
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Pred. No.
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Pred. No. 8.3e+04;
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1998-04-21
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J. 5.7e+04;
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RESULT 14
US-60-427-808-556622
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; TYPE: DNA
; ORGANISM: Mus musculus
US-60-427-808-320640
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SOFTWARE: Microarray Probe Sequen
SEQ ID NO 320640
LENGTH: ACCOUNTS AND ACCOUNTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
CURRENT APPLICATION NUMBER: US/60/427,808 CURRENT FILING DATE: 2002-11-20
                                                                                APPLICANT: Xue Mei Zhou
TITLE OF INVENTION: Met
FILE REFERENCE: 3528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 320640, Application US/60427808 GENERAL INFORMATION:
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 875
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CURRENT FILING DATE: 2002-11-20
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TITLE OF INVENTION: Methods of Genetic Analysis of Mouse
FILE REFERENCE: 3528
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IIOR, FILING DATE: 2001-05:11
IIOR APPLICATION NUMBER: US 60/291,311
IIOR FILING DATE: 2001-05-17
IIOR PPLICATION NUMBER: US 60/353,058
IIOR APPLICATION NUMBER: US 60/361,293
IIOR APPLICATION NUMBER: US 60/361,293
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US-60-427-808-917221
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SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 556622
                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION
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                                                                                                                                                                                                                                                                                     FILE REFERENCE: 3528
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                                                                                                                                                                            TYPE: DNA ORGANISM:
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AUTHORS
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Fax: 314 286 1810
Email: est@watson.wustl.edu
Insert Size: 603
High quality sequence starts: 1
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Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P. and Wilson,R.
The WashU-Merck EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        48 bp mRNA linear EST J yo58c08.sl Soares breast 3NbHBst Homo sapiens cDNA clone IMAGE:182126 3' similar to gb:X06409 RAF PROTO-ONCOGENE - SERINE/THREONINE-PROTEIN KINASE (HUMAN);, mRNA sequence.
                                                                                                                      Unpublished (1995)
Contact: Wilson RK
Washington University School of Medicine
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                         H26959
                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AU2770282 AU2770282
AZ310534 TM0025503
AL587842 AL587842
AZ512567 1M0358017
A1036793 Homo Sapi
AZ602556 IM0421E20
A1267816 a935099 x
AU102743 AU102743
BEF013308 rk29202.y
AA737998 nx15b01.s
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AZ774766 2M0004G05
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REFERENCE
AUTHORS
     FEATURES
                                                                                                                                                                                                                                                                                                                                                                                                                                     SOURCE
ORGANISM
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VERSION
KEYWORDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AZ425831/c
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Best Local
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                                                                                                                                                                                                                                                        JOURNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 TTTGGGTTTNAATTNNGTTTTNATGGCACCGGACAAATACAGTTATC 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TTTGGGTTTTAATTTTGTTTTTATTGCACCTGACAAAATACAGTTATC 2932
                                                                                                                                                                                                                                                                              and Wright, D., Weiss, R. Mouse whole genome sca
                                                                                                                                                                                                                                                                                                                                                                               Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GSS.
                                   Plate: 0206 row: D column: 14
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                              Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AZ425831 47 bp DNA linear GSS 0: 1M0206D14F Mouse 10kb plasmid UUGCIM library Mus musculus clone UUGCIM0206D14 F, DNA sequence.
                                                                                  Pel: 801 585 5606
Pax: 801 585 7177
Pamail: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
                                                                                                                                                                                                University of Utah
University of Utah
                                                                                                                                                                                                                                     Contact: Robert B.
                                                                                                                                                                                                                                                    Unpublished (2000)
                                                                                                                                                                                                                                                                         plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                             Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AZ425831.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AZ42583
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IMAGE Consortium (info@image.llnl.gov) for Trace considered overall poor quality Insert Length: 603 Std Error: 0.00.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    house mouse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This clone is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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Seq primer: Promega ~21m13
                                                                                                                                                                                 .308,
quality sequence stop: 47
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12
                                                                                                                                                         ASD
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                                                                                                                                                                           Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          double-stranded CDNA was ligated to Eco. RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT7T3 vector (Pharmacia). Library went through one round of normalization to a Cot = 20. Library constructed by Bento Soares and M.Fatima Bonaldo."

9 g 17 t 4 others
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        'lab_host="DH10B (ampicillin resistant)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 'sex="Female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              'clone_lib="Soares breast 3NbHBst"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /db_xref="taxon:9606"
/clone="IMAGE:182126"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             'dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    'db_xref="GDB:3815
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GI:10549940
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                                                                                                                                                                                                                  Genome Center
                                                                                                                                                                                                                                                                                    scaffolding with paired
                                                                                                                                                                                                                                     Weiss
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Pred. No. 7
                                                                                       Std Error:
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ORIGIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                    Clone from S. Wiemann, sequenced by sequencing consortium of the German
                               Berlin-Charlottenburg,
                                                This clone is available at the RZPD in Berlin Please contact the RZPD: Ressourcenzentrum, H
                                                                                                                                                           Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases. MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2784 GGCCAGCCTGTTTCTCTAACAAACAAACAAACAAACAACAGC
                                                                                       No s1 sequence available
                                                                                                                                                                                                                                                                                                              Homo sapiens (human) Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                12-MAR-1999
12-MAR-1999
                                                                                                                                                                                                                 Bloecker H., Boecher M., Brandt P., Mewes W., Gassenhuber J.,
                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                      EST; expressed sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AL037866.1
                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens mRNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AL037866;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HSM002199
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       47 GCCCAGGTTATTTTCAGAACAAACAAACAAACAACAAC 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             , 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard; RNA; EST; 45
                                                                                                                                                                                                                                                                                                                                                                                                                                              (Rel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."

1 2 24 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid Rl. The vector was ligated with adaptors complementary to the insert adaptors and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (http:/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone_lib="Mouse 10kb plasmid UUGC1M library
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone="UUGC1M0206D14"
                                                                                                                                                                                                                                                                                                                                                                                                                                                59,
                                                                                                                                                                                                                                                                                                                                                                                                         EST DKFZp564J147_r1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Created)
Last updated,
                               GERMANY;
                                                                                                                                                                                                                                                                                                                                                                      tag.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 24.6; DB 17
Pred. No. 1.8e+05;
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                                                                                                      GBF within the cDNA Genome Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                Version 1)
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                                              Heubnerweg, 6, 14059
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2822
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                                                                                                                                                                                                                                                                                                                Euteleostomi; Mammalia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                     Wiemann
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AZ391073
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SOURCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
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                                                                                                                                                                                                                                                                                                                              High quality sequence stop: 40 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and Wright, D., Weiss, R.
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                                                                                                                                                                                                                                                                                                                                                                                       Plate: 0152 row: I column: 24
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                        Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Err
Plate: 0152 row: I column: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       University of Utah University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  plasmid
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801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                       plasmid ends
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone_llb="564 (synonym: hfbr2). Vectcr pAMP1; host
X1-2blue; sites NotI + SalI"
/dev_stage="fetal"
           (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/clone="DKFZp564J147"
                                                                                                                                          /lab_host="E. Coli strain XL10-Gold, Tl-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                            Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                       /db_xref="taxon:10090"
/clone="UUGC1M0152I24"
                                                                                                                                                                                                                                                                            /organism="Mus musculus"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /tissue_type="brain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /db_xref="taxon:9606"
adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                      /sex="Male"
                                                                                                                                                                                                                     /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%;
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Pred. No. 5.4e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                            Std Error:
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2872 GTTTTCTTGATGTTTTGGGTTTTAATTTTGTTTTTATT 2909

Matches Query Match

28;

Conservative

0;

Mismatches

Indels

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Gaps

0;

Local

Similarity

0.7%; 73.7%;

Score 22; DB 12; Pred. No. 8.7e+05;

Length 43;

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KEYWORDS
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ORGANISM
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ORIGIN
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2872 GTTTTCTTGATGATTTGGGTTTTAATTTTGTTTTTATT 2909
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov
Tissue Procurement: CLONETECH Laboratories, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BE969098.1 GI:10581742
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (bases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and
                    (where B = A, C, or G and N = A, C, G, or T). Average insert size 1.25 kb (range 0.6-4.0 kb). 14/15 colonies contained inserts by PCR. This library was enriched for full-length clones and was constructed by Clontech Laboratories (Palo Alto, CA)."
                                                                                                                                                                                                                                                               /Clone_IIAGE:333811"
/Clone_Lib="NIH_MGC_74"
/Clone_Lib="NIH_MGC_74"
/Clone_Lib="NIH_MGC_74"
/Clone_Lib="NIH_MGC_74"
/Clone_Tib="DH10B (T1 phage-resistant)"
/Iab_host="DH10B 
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0; Mismatches
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BJ047667/c
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      TITLE
                                                                                    AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                      28.71 GGTTTTCTTGATGATTTTGGGTTTTAATTTTGTTTTTATTTGCAC 2913
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                                                                                                                                                                                                                49 bp uk62a04.y1 Schiller mouse MAC16 Mus IMAGE:1973550 5', mRNA sequence.
    Waterston, R.
The WashU-HHMI Mouse
                                                                                                                                                                                EST
                                                                                                                                                                                                                                                                                                                                                                                                     29;
                                                                                                                                                                                                              AI316598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BJ047667 NIBB Mochii normalized Xenopus neurula
BJ047667 NIBB Mochii normalized Xenopus neurula
                                                                                                                  Mammalia;
                                                                                                                               Eukaryota; Metazoa;
                                                                                                                                               Mus musculus
                                                                                                                                                            house mouse.
                                                                                                                                                                                             AI316598.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           National Institute of Genetics
1111 Yata, Mishima, Shizuoka 411-8540, Japan
Tel: 81-559-81-6856
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: tshini@genes.nig.ac.jp.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Expressed genes in X. laevis embryo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Xenopodinae; Xenopus.

1 (bases 1 to 47)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            African clawed frog
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Tadasu Shin-i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kitayama,A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Xenopus laevis
                                                                                                (bases 1 to 49)
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                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              81-559-81-6855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note-"Vector: pBSRN3; Site_1: NotI; Site_2: EcoRI; cDNAs were oligo-dT primed and directionally cloned. Staging according to Nieuwkoop and Faber. Library is subtracted and was constructed by N. Garrett and A.M. Zorn, (Wellcome/CRC Institute). "
                                                                                                               Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (Wellcome/CRC Institute). "
4 c 3 g 3 t
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/clone="XL017e20"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      dev_stage="stage 15"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'tissue_type="whole_embryo"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               organism="Xenopus laevis"
                                                                                                                                                                                              GI:4031865
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                                                                                                                                                                                                                                                                                                                                                                                                                0.7%;
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                                                                                                               Chordata;
Rodentia;
   EST Project
                                                                                                                                                                                                                                                                                                                                                                                              Score 21.8; DB 13;
Pred. No. 1e+06;
0; Mismatches 14;
                                                                                                          Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                             Bowles, M., Dietrich, N., Dubuque, T.,
                                                                                                                                                                                                                                                       mRNA
                                                                                                                                                                                                                                        musculus
                                                                Martin, J., Morris, M.,
                                                                                                                                                                                                                                                                                                                                                                                              Indels
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                                                                                                                                                                                                                                                       linear
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                                   Wilson, R: and
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library Xenopus
                                                                                                                                                                                                                                                     EST 17-DEC-1998
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; Pipidae;
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AUTHORS
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AZ593107/c
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Best Local
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                                                           Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                          University of Utah University of Utah
                                                                                                                                                         Unpublished (2000)
Contact: Robert B.
                                                                                                                                                                                                                   and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                               1 (bases 1 to 49)
Dunn, D., Aoyagi, A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  32; Conservative
                                                                                                                                                                                        plasmid inserts
                                                                                                                                                                                                                                                                                                                               Mus musculus
                                                                                                                                                                                                                                                                                                                                                                              AZ593107.1
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Seq primer:
                                                                                                                                                                                                                                                                                                  Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                 house mouse.
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                                                                                                            308,
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                                                                                          USA
                                                                                                            Biomedical
 CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                              GI:11715297
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JOURNAL
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This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WashU-HHMI Mouse EST Project
Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. La
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seq primer: -40RP from Gibco
Location/Qualifiers
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                                                                                                                                                                                                                                 /note-"Vector: pBluescript SK- (Stratagene); Site_1: Ecc. Site_2: XhoI; Double-stranded cDNA was prepared from cell-line MAC16 using primer 5'-GAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGTCTGAGT(18)-3'. An EcoRI adaptor was used on the 5' end of the cDNA as follows: 5'-AATTCGGCACGAG-3'. The library was size-selected and went through one round of amplification. Average insert size is 1.7 kb, with a range from 0.4-12 kb. This library was size is 1.7 kb, with a range from 0.4-12 kb. This library was size is 1.7 kb, with a range from 0.4-12 kb. This library was size is 1.7 kb, with a range from 0.4-12 kb. This library was size is 1.7 kb, with a range from 0.4-12 kb. This library was size is 1.7 kb, with a range from 0.4-12 kb. This library was size is 1.7 kb, with a range from 0.4-12 kb. This library was size is 1.7 kb, with a range from 0.4-12 kb. This library was size is 1.7 kb, with a range from 0.4-12 kb. This library was size is 1.7 kb.
                                                                                                                             University). "
                                                                                                                                                                                                    was constructed by Dr. Martin Schiller (Johns Hopkins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone_lib="Schiller mouse MAC16"
/cell_line="colon cancer cell line"
/lab_host="SOLR"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone="IMAGE:1973550"
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/db_xref="taxon:10090"
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kb. This library
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2872 GTTTTCTTGATGATTTGGGTTTTAATTTTGTTTTTATTGCACCTGACAA 2920 Score 21.8; DB Pred. No. le+06;); Mismatches 17; Indels 0 Gaps 0

Email: ddunn@genetics.utah.edu Insert Length: 10000 Std Error: Plate: 0404 row: O column: 22 1M0404022F Mouse 10kb plasmid UUGC1M library clone UUGC1M04040402 F, DNA sequence. Mouse whole genome scaffolding with paired Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. Genome Center Weiss Chordata; Rodentia; Polymers Research Bldg., Barber, M., 49 bp Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus Beacorn, T., 0.00 end Duval,B., Mus 20 reads Š musculus 2030 GSS 13-DEC-2000 from 10kb Hamil, [H] Reilly SLC,

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RESULT 9
HSM003616
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ORIGIN
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                                      Clone from S. Wiemann, sequenced by sequencing consortium of the German No s1 sequence available
                                                                                                                    Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2803 CAAACAAACAAACAACAGCCTTGTTTCTCTAGTCACATCATGTGTATA 2851
                                                                                                                                                                                                                                                                                                                                     EST; expressed sequence
                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens mRNA;
                                                                                                                                                                                                                                                                                                                                                                                                                     12-MAR-1999
12-MAR-1999
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  This clone is available at the RZPD in Berli Please contact the RZPD: Ressourcenzentrum,
                                                                                                                                                                              Bloecker H., Boecher M., Brandt P.,
                                                                                                                                                                                                                                                                                                Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         49 CACACACACACACACAGAGTTTTGCATGAAGTTATGTAATGTGTACA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Class: plasmid ends
High quality sequen
                                                                                                                  Klopferspitz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard; RNA;
                                                                                                                                                                                                                                                                                                                                                                                                               (Rel. 59, Created)
(Rel. 59, Last updated, Version 1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."

8 c 11 g 18 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWDA2 (gil4732114|gb|AF729072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             musculus C57BL/6J (male) was obtained
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /db_xref="taxon:10090"
/clone="UUGC1M0404022"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Mus musculus"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note-"Vector:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           'clone_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                                                                                                                                                                                                                                                                                                                              EST DKFZp566L194_r1
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                                                                                                                  Martinsried, GERMANY
                                                                                                                                                                          Mewes W., Gassenhuber J.,
                                                                             GBF within the
                                                           Genome Project
                        Berlin
                                                                                                                                                                                                                                                                                                                                                                              (from clone DKFZp566L194)
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Heubnerweg 6, 14059
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                   source
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Takeuchi,I., Kohara,Y. and Tanaka,Y.
Population analysis of cDNAs from unicellular and multicellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AU270282 VS Dictyostelium discoideum cDNA clone VSJ664 3', mRNA
                                                                                                                                                                                                                                                                                                                                                        Contact: Hideko Urushihara
Institute of Biological Sciences
                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Mycetozoa; Dictyosteliida; Dictyostelium
1 (bases 1 to 50)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dictyostelium discoideum.
                                                                                                                                                                                                                                                                                         Email: hideko@biol.tsukuba.ac.jp
                                                                                                                                                                                                                                                                                                                               l-1-1 Tennoudai, Tsukuba, Ibaraki 305-8572, Japan
                                                                                                                                                                                                                                                                                                                                           University of Tsukuba
                                                                                                             Similarity
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X1-2blue; sites NotI + SalI"
/dev_stage="fetal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/clone="DKFZp566L194"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                             /dev_stage="vegetative"
3 c 1 g
                                                                                                                                                                                                                 /clone="VSJ664"
                                                                                                                                                                                                                              /strain="AX4"
/db_xref="taxon:44689"
                                                                                                                                                                                                                                                                             Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /db_xref="taxon:9606"
                                                                                                                                                                                                     /clone_lib="VS"
                                                                                                                                                                                                                                                    organism="Dictyostelium discoideum"
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Pred. No. 1.1e+06;
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                                                                                                              Score 21.4; DB 9;
Pred. No. 1.3e+06;
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                                                                                                   Mismatches
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                                                                                                                           Length 50;
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DEFINITION LOCUS

AZ310534 43 bp DNA linear GSS 29-SEP-2011M0025J03R Mouse 10kb plasmid UUGCIM library Mus musculus genomic

29-SEP-2000

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RESULT 12
AL587842
LOCUS
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        DEFINITION
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           JOURNAL
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                                                                                                                                                                        2801 AACAAACAAACAAACAACAG
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                                                                                                                             5 AACAAACAAACAAACAAACAG
      AL587842
AL587842 BP Chicken
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0025 row: J column: 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 High quality sequence stop: 43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse whole genome scaffolding with paired end reads from 10kb
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AZ310534.1 GI:10352617
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (bases 1 to 43)
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                                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                                                                                                                              10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|aF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Biomedical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /db_xref="taxon:10090"
/clone="UUGC1M0025J03"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Mus musculus
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /lab_host="E. Coli strain XL10-Gold, T1-resistant,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                                                                                                                                                                                 0.7%;
44 bp mRNA linear I
Brain Library Gallus gallus cDNA
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Rodentia;
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                                                                                                                                                                                                                                 Score 21;
Pred. No.
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Sciurognathi; Muridae;
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1.6e+06;
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                 University of Utah Genome Center University of Utah
                                                                                                                     Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                     Eukaryota; Metazoa;
Mammalia; Eutheria;
1 (bases 1 to 36)
                                                                                                                                                                                                                                                                                                               36 bp DNA linear GSS 05-OCT-20010358G17F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0358G17 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27;
                                                     Contact: Robert B.
                                                                   Unpublished (2000)
                                                                                                     Mouse whole genome
                                                                                                                                                                        Dunn, D., Aoyagi, A., Barber, M.,
                                                                                                                                                                                                                                            Mus musculus
                                                                                       plasmid inserts
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Gallus gallus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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Biomedical Polymers Research Bldg.,
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0 c 4 g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /tissue_type="Brain"
/dev_stage="Unknown"
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73.08;
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Rodentia;
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Pred. No. 1.6e+06;
0; Mismatches 10
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 SLC,
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Email:

ddunn@genetics.utah.edu

Insert Length: 10000

Std Error: 0.00

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RESULT 14
HSM001110
ID HSM000
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XY AL036
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Best Local Similarity
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Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens mRNA; EST DKFZp564K0763_r1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AL036793.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2797 CTCTAACAAACAAACAAACAAACA 2820
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AL036793;
                                                                                                                                                                                                                                                                                                    Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HSM001110
                                                                    Duesterhoeft A., Lauber J., Mewes W., Gassenhuber J., Wiemann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             L2-MAR-1999
L2-MAR-1999
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                                                                                                                                                                                                                                                                                                                                                                     expressed sequence tag
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    High quality sequence stop:
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Seq primer: CGTTGTAAAACGACGGCCAGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard; RNA; EST; 44 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (Rel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwp42 (gi14732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and selected for ampicillin resistance.
0 c 15 g 21 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (http://www.jax.org/resources/documents/dnares/).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note="Vector:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /sex≂"Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     'clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /db_xref="taxon:10090"
/clone="UUGC1M0358G17"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       organism="Mus musculus"/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     59,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.7%;
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_host="E. Coli strain XL10-Gold, T1-resistant,
_ host="E. Coli strain XL10-Gold, T1-resistant,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Created)
Last updated, Version 1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 20.8;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               coli XL10-Gold (Stratagene) cells
                                                                                                                                                                                                                                                                                                                                                                                                                                             (from clone DKFZp564K0763)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .DB 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17;
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KEYWORDS
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AZ602556/c
                                                                                                         FEATURES
                                                                                                                                                                                                                                                                                                              COMMENT
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Best Local (
                                                                                                                                                                                                                                                                                                                            JOURNAL
                                                                                                                                                                                                                                                                                                                                                           TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2772 GGCGGGCCGCAGGCCTGTTTCTCTAACAAACAAACAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Clone
                                                                                           source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 44 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This clone is available at the RZPD in Berlin Please contact the RZPD: Ressourcenzentrum, Heubnerweg Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequencing consortium of the German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MIPS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         No s1 sequence available
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Am Klopferspitz 18a D-82152 Martinsried, GERMANY
                                                                                                                                                                                                                                                                                                                                                                                   Dunn,D., Aoyagi,A., Bar
Islam,H., Longacre,S.,
M., Rose,M., Rose,R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AZ602556 45 bp DNA linear GSS 13 1M0421E20F Mouse 10kb plasmid UUGCLM library Mus musculus clone UUGCLM0421E20 F, DNA sequence.
                                                                                                      High quality sequence stop: 45.
Location/Qualifiers
                                                                                                                                                   Plate: 0421 row: E column: 20 Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                 Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
                                                                                                                                                                                                                   Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                              University of Utah University of Utah
                                                                                                                                                                                                                                                                                                                         Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                           plasmid inserts
                                                                                                                                                                                                                                                                                                                                                        Mouse whole genome scaffolding with paired
                                                                                                                                                                                                                                                                                                                                                                         and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AZ602556.1
                                                                                                                                                                                                                                                                                                            Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   house mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                        lass: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                   (bases 1 to 45)
                                                                                                                                                                                                                                                                 308,
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                                                                                                                                                                                                                                                   USA
                                                                                                                                                                                                                                                        Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21 A; 6 C; 10 G; 4 T; 3 other;
                            /organism="Mus musculus".
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0421E20"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
/clone_lib="Mouse 10kb plasmid UUGC1M library"
/sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      X1-2blue; sites NotI + SalI*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone_lib="564 (synonym: hfbr2).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
/clone="DKFZp564K0763"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /tissue_type="brain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /dev_stage="fetal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GI:11724746
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                                                                                                                                                                                                                                                                                            Genome
                                                                                                                                                                                                                                                                                                            Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                    Barber, M., Beacorn, T., Duval, B., Hamil, C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 20.8;
                                                                                                                                                                                   Std Error:
                                                                                                                                                                                                                                                                                                                                                                                      Stokes, R.,
                                                                                                                                                                                                                                                                                                                                                                                   Mahmoud, M., Meenen, E., Pedersen, T., Reilly Stokes, R., Tingey, A., 'von Niederhausern, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                              Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              No. 1.8e+06
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                                                                                                                                                                                    0.00
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                                                                                                                                                                                                                                                                                                                                            end reads from 10kb
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                                                                                                                                                                                                                                                                20 S. 2030 E., SLC, UT
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/lab_host="E. Coli strain XL10-Gold, T1-resistant,

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BASE COUNT
ORIGIN
Search completed: May 16, 2003, 10:26:05
Job time : 4055 secs
                                                                                                                                                                                                                                                 Query Match 0.7
Best Local Similarity 91.7
Matches 22; Conservative
                                                                                                                     (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWDA2 (gil4732114 [gb]AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli xL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                 Score 20.8; DB 17;
Pred. No. 1.9e+06;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                             Length 45;
                                                                                                                                                                                                                                                 Gaps
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